

5 SEQ ID NO:6132
 SEQUENCE LENGTH:21
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07290
 SEQUENCE DESCRIPTION:
 GATCATAAAT AAATGTTTAA A 21

10 SEQ ID NO:6133
 SEQUENCE LENGTH:331
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS07291
 SEQUENCE DESCRIPTION:
 GATCATTTC AAAAATTGCA TTGGGGCCTT CGTTATTTAC CATAGTATTT NCACTTTCAT 60
 AGTTTTGTCA CCTTTTGTGA CTGTGAACAG TTCAACCAGT GACCGACTTC TCTCTCATGC 120
 TGTTTACCCC ACACACAATT TCCCCTCAAA TTCTGAAAAAT AAGAACCCTGT TAATAGGTTG 180
 20 GAAAGCTGTG TACTCTATTC ATATAATTGC CTTTCATGCT AGTGGAGAGT GGTGTCATTA 240
 GCATCTNAAT TTTAGAGTTG TGNAATGATT TTNCCAATTA GGAATTGANT GTGTATTNT 300
 NTCCTGTTTA ATAAGNNGNG CAAATNTGNT N 331

25 SEQ ID NO:6134
 SEQUENCE LENGTH:40
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07292
 30 SEQUENCE DESCRIPTION:
 GATCTTAAAA CAAATGAAA TAAACGTGAA AAGGAGCAAA 40

35 SEQ ID NO:6135
 SEQUENCE LENGTH:53
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07293
 SEQUENCE DESCRIPTION:
 40 GATCGTGAAT AGTAAGTATC ACTTCCATAA ATAGAAAGTG TCAGAAAAAT AAA 53

45 SEQ ID NO:6136
 SEQUENCE LENGTH:336
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07294
 SEQUENCE DESCRIPTION:
 GATCTTGAAA GAGGTAAAAT AAACCTTCAC TGCTCCACTC CAGGTGAATC CGCCCACTCC 60
 50 CACTGACCTA GTAGAATTG TAATTTAATA CTTACCTTCT ATTTCTGAAA TCAGTTGTGA 120

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ACTGTTGCCT TATGTTTACA GGTTTAAGAA CCTCAGTGAA TTCATTTTTT AAAATCTGCT 180
 ATTCTGAGAA GCATTGAATG AATTCTTAAC AAGAAGNCTC ATCTGTAGCT GTTTGCTGAC 240
 TCCTATGAGC CCATAAGGGT TCTGTGCTTA GCATTACAA ANTAAGGTTT ATAGGGTAAA 300
 AGCCAATGTA TTANTTTTTT TTTGCATGGG GGGCNN 336

SEQ ID NO:6137
 SEQUENCE LENGTH:235
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07295
 SEQUENCE DESCRIPTION:
 GATCTAGGAC ACTCATATCC AGGGTGATTC TCTCATAAG GAGGCATGGG CTGAGTCTCC 60
 ATTTATAGGG GCTGTCCCA TAGAGACTTC AAGCCACACC TTTCTAGGGA TTCAGGTTGC 120
 ACATTACCTG ATTCTCACAG TTGATTTGAG CCTGCAAGAC TGTAACCCA TCCCATTATC 180
 ATGTTTTCTT CCTCATTAC CTACATTAT CATTAAACCT TTGTGATTCA GCAA 235

SEQ ID NO:6138
 SEQUENCE LENGTH:142
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07296
 SEQUENCE DESCRIPTION:
 GATCCAGGAT GACTGTACAT CTCAGATAAA GGAGGCTCAG CGCTGGAAGG ACAGCTGGAA 60
 GCAGTCCCTG CACACTATCC AGGGCCTGTA TGTGTGACCC TCCGCCCCAC CATGAATAAA 120
 CACTTTCTTA TACAGACTCA AA 142

SEQ ID NO:6139
 SEQUENCE LENGTH:277
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07297
 SEQUENCE DESCRIPTION:
 GATCCGCCAA GCAGGGCAGT ATCTGGGTTA CAGCCACAGC AGGAAGAACC ACTGGAGAGA 60
 CACTCAGAAG GACTTCCCTG ACATCTGCAC CAAGGAGAGT CTGGACAAAG CTAAGTAAAC 120
 TCCTCCACCT GCTTCTAAAA TGCCAGAAG CCAACTGGAC CNGTTTGTA TTTNGAAGTA 180
 GGNGGNTGGN GGCTTTAACC NCCAAAGGGT TCCTGAAGCC CNGGAACCTG GANTNTGGAG 240
 GGGGTTNNGG CCTTANTTGA GGGTTGGTGG GTTTCNN 277

SEQ ID NO:6140
 SEQUENCE LENGTH:247
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07298
 SEQUENCE DESCRIPTION:
 GATCAGAAAA CATTGATTTA TATTTAAAAA TAGGATATTG GAGCCATTGT CTTAGATAAT 60

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TNNCAAGGCA AGANCAAGCC TGCAATAAAT AATGAGACTT TGACATTCTT TGGCGAAGAA 120
 GATGTCTCCT GCCCTAATAT GACTGTCACT CAGCAATTC ANTTTTNTGT TTACTTGATT 180
 TANAANGGGA GGATAANTTG TCCTGANAAG TTTTTTGTG TCCTTAACAT TTCANNGTTT 240
 TCNCTGN 247

SEQ ID NO:6141
 SEQUENCE LENGTH:228
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07299
 SEQUENCE DESCRIPTION:
 GATCCCTGAT GGGATGCCTG AAACCCAAAA TACTCTTAAG TCTTGATTTC TAGGGCCTTC 60
 TAAGGCAAAA NATTGAGAGA CCAAAAAAGA GAAAATATGT TTAGTAAAGA AAATATTTNC 120
 TTGACTAGT TATGATGGTC ATATAGGGGA GANAAGGCAA ATTTTGGNAA TGTAAGATGG 180
 GGACANTTAA TGTNTATNCN CTGTCCAANT TCAANAGGTA CAACNGAN 228

SEQ ID NO:6142
 SEQUENCE LENGTH:228
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07300
 SEQUENCE DESCRIPTION:
 GATCCACCTG TCTCTTGAAA ATATAAAAAG AAAAAACAGG TTGCGCTTGG CATCAGAGAG 60
 CACAAAGATT AAAAGTTACT TTAAATTTGC CAATATTTTG GGAGANCAAT AAANCTACAT 120
 TTTTCCTCT TCCATACTGG TAGATGCGAA ATTNATCTGT GCATGANAGG GTCACCTCTG 180
 TNATCGTGCA ACAGNTTGG TATTANNCCT TNNCTGTGGT TTTNCCAN 228

SEQ ID NO:6143
 SEQUENCE LENGTH:299
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07301
 SEQUENCE DESCRIPTION:
 GATCCTCTGA TATACAATTA GAGATATTTT AATATAGNCC CCAAGCATTG TGTGCATAAA 60
 AGTTAACATT AGGCTGTGGT GCAGTAACCA TTAAATGTCG AGGCTCTATT TCGGAAATAC 120
 ACTACAAATG TTAAGTACG TGGCTGTCCT CTTAAGNCAC TAGTAGAGCA AAGCCTTAAT 180
 CATATCANCT TAATTCTGTT ACACAAATATG TGTTTTTNAA TATACTANCC ATTTCTNATG 240
 GAANGGTCCT GTGGGGNGCC CNTCCTCTCG CCANGCCATC ACAGGCTCTG CATAACGN 299

SEQ ID NO:6144
 SEQUENCE LENGTH:152
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07302
 SEQUENCE DESCRIPTION:

GATCGGGTGC TAGGACAAC TGCAGTCCAAT CCACCAGCTC TCCNTGCCCC TGTGTCTTAT 60
 TTCAGACATG AGAATAACTG TACAGTGTA ACTTATAAAG CGTTTTTAAT GGTGTAGAT 120
 TGGAAATAAA GTATGTCATA TGAACAGCTA AA 152

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SEQ ID NO:6145
 SEQUENCE LENGTH:131
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07303

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SEQUENCE DESCRIPTION:

GATCCACCT GCTCAGGCCG TACACCTTAT GGGGAGCCAT TAAGGGGNC CATCTAGGCA 60
 TTCTGGNTAC AGCCAGTGC TCATCCAGC GTATGCTCCN CTCNTAGNG CAGCCTAAAG 120
 NGTCAGCCAC N 131

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SEQ ID NO:6146
 SEQUENCE LENGTH:154
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07304

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SEQUENCE DESCRIPTION:

GATCTTCAGA GAAATAAAAC TTGAGTGCAG TAGTTTATTT GCGGTTTGGC TCCAGGATGC 60
 AGGNGTGGGA AAGATGGTAG AGAGCCAGAA GAAAANTCAA TAGAACGGTA TTACGGCGNN 120
 TGGNTATATA ATNCTTCCCN GAATTGTCTG ACTN 154

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SEQ ID NO:6147
 SEQUENCE LENGTH:251
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07305

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SEQUENCE DESCRIPTION:

GATCCAGTGC TATTCTGTCA CCAACCTAAG AATCCCAATT GCACCTTCTG TTTCTGACAG 60
 TCACAGGTGA CAGCTGNGGN TCTATAATAC AGACTGGTGT CTTAGAGGTA GGAATAATAC 120
 ATGATTATNA AGCATCACCC TGCTAATACA TAATAATGTC TTTTATATT ATAAGNGATT 180
 GAGTTTAGTT CATTNCAATA CATTGTACN TGTA AAAANT GCAANGTNGA CCTNNGTAAT 240
 CNTTTAATCG N 251

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SEQ ID NO:6148
 SEQUENCE LENGTH:213
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07306

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SEQUENCE DESCRIPTION:

GATCTCGGAG ACTCNGTCCA TTCTAAGATA CCATGTTTCC ATGACCCTAC ACACAATAGC 60
 TACACAGGTG GAGCAACATG AGTCTGAGNA GCTTGCACA AAAGANGCTT GTTTTCCACA 120
 AATGNANCTG GGGTGGTTAA CTTCTAATTG TACCCTGTCT TAGCGTTCAG ATGCATTGGT 180

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TCCTTCCCCA ATGTGACCAT CTNNNNATTC NAN

213

SEQ ID NO:6149

SEQUENCE LENGTH:113

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07307

SEQUENCE DESCRIPTION:

GATCAAGATG AATNTAATGT ATAATCAGAA TCCTAGGTAA GTAAACAAAA AAAGGGTTTT 60
 TTTNAGAATC AAACCTTTGG GGATTGGGTG GTAAAGGGAA ATATTNTNT NCN 113

SEQ ID NO:6150

SEQUENCE LENGTH:228

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07308

SEQUENCE DESCRIPTION:

GATCTGTGCA TCTGGCCAGC ACTGTGGTGA CATCCTTAGG AATCCATGGG GAGAGAGAAA 60
 GCATTCAGGA GTTAGTGGGT CACGTTTGAC AAGGGCCAAT AAAGAAATAT GCAAAGACAA 120
 AAANCAAGAA GNACATTGTC ATATTNCTA CCTTTTGTTC ATATAAATTT ATNTCANTGA 180
 TTTTCGCTTN TGTTAATNTG CAATGTATAT NNTATGCTAA CNTTTNCN 228

SEQ ID NO:6151

SEQUENCE LENGTH:42

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07309

SEQUENCE DESCRIPTION:

GATCACAATT GTGGACATTA AAACAGAAAC TGTTACACAA AA 42

SEQ ID NO:6152

SEQUENCE LENGTH:311

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07311

SEQUENCE DESCRIPTION:

GATCCAGCTT ATNCTTTAAT TTTCAAGTCC ATTCTTGGGG CTGGTGGGGA GGCAGGAGAA 60
 TACCCCTCCC TAAGCCCTTA GTGTGTGCCG AGCTTGCTTT GTNATGTTGG CAGGGGAGGG 120
 GAGACCTGGG TGGTGA CTGA GTTCCCTTTA TCAAACCCTT CAATGGGCAC AAAATTGAGT 180
 GCTTGATTTT AGGTTTTATT TTTTATGAA TGTCCAAATC TGTGTTTCCC CCTGCCATCC 240
 CAGACTNTGT GGCCAGTTNA AAAGTGTCTG GNNTGTGTTT ANCTCTNCCC TAANATNCTG 300
 GAGNAGGGGT N 311

SEQ ID NO:6153

SEQUENCE LENGTH:270

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07312

SEQUENCE DESCRIPTION:

GATCTNNGTG GGTAGCCTCA AGCTGGAGAC ATTGTCTGNC CTCACTGCCC ACTGGGTAGA 60
GACACCTGGA TGACATTTC GTACCCGCTG GCAGCTCCTC CCACTCATCC AGAGACGGCC 120
GGTGGGCAGG CGGGGAGAGC AGCTTTCCTT CCCTGGNGAG AACCGTCTTT ACCTCAGCTC 180
CTAGGGNCTC CAGCCAACAA AGCTNCAGCA CGGTGACTNG GTGATGGGNG ACCAATNNCA 240
GAAGTAAANN GACTGTAGAC CNATATTCCN 270

SEQ ID NO:6154

SEQUENCE LENGTH:40

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07313

SEQUENCE DESCRIPTION:

GATCAAAATC TAAATAAAG TTGCCTGTTG TGACTTTAAA 40

SEQ ID NO:6155

SEQUENCE LENGTH:130

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07315

SEQUENCE DESCRIPTION:

GATCTATACT TAACATATTT AAAATAGTGA TGATTGAAAT AACTGCAAGG TATAGATGNA 60
ACTGAAGGNT TTCACATTCC TTTTTGTTT ATCATTGGA ATTNCTGTA TTTANCTGNC 120
TAGAGANCAN 130

SEQ ID NO:6156

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07316

SEQUENCE DESCRIPTION:

GATCATATCA CTCAATGTGG CCATTAATAT TTCTGGAGAA GACACAAACC TATTATAAT 60
ACTTTTAAAA AATCCGAAGT GATAATAAGA NTGTTTTTAC ATTGAAAGTA ATACTTAAGT 120
GTTTACATAA CATTCTGTAT GCACTCAAGA AAATAAAGTT TTGTTAATAC AAA 173

SEQ ID NO:6157

SEQUENCE LENGTH:248

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07317

SEQUENCE DESCRIPTION:

GATCTCAGTT ACAGTTTTT TTTCTTTTT AATTCATTA TTTGGGTTT TTGGTTTTTG 60

5 CAGTCCTATT TATCTGCAGT CGTATTAAGT CCTATTGCTA GAATAGGTTA CTNNANGAAA 120
 GATTATATTC TGAAAGAAAA ATAACGTGACA TTATATATAA CCAATTAATT TAANGTATTG 180
 CCATTTAAAT TACACACTGN GAGCATGTCC TATGCAGACA ATAGATTTTN CCTGTNCAAN 240
 NNNTTNTN 248

10 SEQ ID NO:6158
 SEQUENCE LENGTH:187
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07318
 SEQUENCE DESCRIPTION:
 15 GATCCAGCAG CGCAGGCAGG AGCTGGAGCA GTCGCTGGTT GTNCGCAACA CCTAGGAGCC 60
 CAAAAATAAG CAGCAGGACG GAACTTTCAG CCGTGTCCCG GGCCCCAGCA TTTNCCCCG 120
 GGCTCCAGCA TCACTCCTCT GCCACCTTGG GGTGTGGGGC TGGATTAATAA GTCATTATC 180
 TGACAAA 187

20 SEQ ID NO:6159
 SEQUENCE LENGTH:21
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07319
 25 SEQUENCE DESCRIPTION:
 GATCATAAAT CCCTGTCTAA A 21

30 SEQ ID NO:6160
 SEQUENCE LENGTH:122
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07320
 SEQUENCE DESCRIPTION:
 35 GATCAAACGT GCTTAAGAGC TAACTCGTGA CACTATGCAG TATTGTTTGA AGACCTGTTG 60
 TTCAACCTCT GTCTCTTTAT GTTAACTGGA TTTCTGCATT AAATGACTGC CCCCTTGTTA 120
 AA 122

40 SEQ ID NO:6161
 SEQUENCE LENGTH:125
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07321
 45 SEQUENCE DESCRIPTION:
 GATCTCTGAC TCATTGTGGC ATATGAGGTG TAAATNTTG TCAAAAAAGA ATCTGGTGTG 60
 TAAACCAAAA TCACACTATG GTGGCATTAA ACCGTTCACT TTTATTCTAA ACNNNANCCA 120
 NCGCN 125

50 SEQ ID NO:6162

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SEQUENCE LENGTH:41

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07322

SEQUENCE DESCRIPTION:

GATCTGTCCA CGACATGGAA AATAAACTGG ATTTTCAGAA A 41

SEQ ID NO:6163

SEQUENCE LENGTH:137

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07323

SEQUENCE DESCRIPTION:

GATCCTATAC GTGAAGGAAC ATATTTAATT TCCTCNCTTT ATATTTCTCG GTTAAAAATAT 60

CGTCATTATA GTTAGCAATT TGAATCTGG CTTACATTGG TTGATACAAA TAANTAATAG 120

ANTANNGCAA NATCAGN 137

SEQ ID NO:6164

SEQUENCE LENGTH:88

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07324

SEQUENCE DESCRIPTION:

GATCTTTTT CTATGCGTAA TCAGACATAC ATATATACTG CAGTGATCT CACGTATTAA 60

TTTTAAAAA TCTTTGTTT TACTTAAA 88

SEQ ID NO:6165

SEQUENCE LENGTH:274

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07325

SEQUENCE DESCRIPTION:

GATCTCTGTA TTGTNTTC AACTAACAAC TAATTNAAA TGTATTGTAT CTTAATNAC 60

CTGTNCTCTT AGATTGINTT TTGCTAGTAA CCCTTACTCA GCTTCTGCCT TTGGGGCTTG 120

GACTAATATT GCTTGTATGG AACTACAGTA CTGTGACTAA ACATGGAGCT CAATGCAGCT 180

ACTGCTTACA ACTGCTTAAA CTTTGTAGTG TGGACTTCAT TTTTCTGT AATAACANCT 240

TATTAAATCC TNGGTGGTAT GCGGGTCCTG GAAA 274

SEQ ID NO:6166

SEQUENCE LENGTH:310

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07326

SEQUENCE DESCRIPTION:

GATCATGTTT AGAAAGGGAC TCATTGGTC CCACTTTGTA ACATGGGAAG AGTAGAAGAG 60

AAAATTAAAT GTGGAAATCA TAAATGCTTT TCTAACAGGG TTATGCTATT CTGTAACACT 120
 AAATGTCTGT CTTTCTCTT AAAAGATTAA GTTTTATTAA TTTTGCATA CATTCTCTA 180
 TTTGANTATA TCTTAACCAT TTTATGTTCC AAATTCGTTT TTGTCCAAG ACATTCTATC 240
 AGANTTTTNC TTGCCTCTAT AAATCTGAAA ANTGGGTTCT AGAATGTCCC ACTTGCTGTC 300
 TCTNAGAGTN 310

SEQ ID NO:6167

SEQUENCE LENGTH:307

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07327

SEQUENCE DESCRIPTION:

GATCCGCCCA CCTCAGCCTC CCAAAGNGCT GGGATNACAG GCATGAGCCA CTGCACCTGG 60
 CCTTAACCCC TCTTTAGATT GGAAAAATA NTTACAACCT TAAAAATAGC TTAGTGTGTA 120
 ACCCTTTGGT AAATAAAGA CCCTTTTATA ATGCACATAT TCCCAACANG GTTAATATAT 180
 TTNGTGAGAT TAAACAAGGC TTGTATATGC TTGACCTTTC TTAANNATG TCCATGTCAT 240
 ACTATTATGA NTGTACATT TTATGAGTCA TNANTATTNT TTTCAAAANG CGCTNCAGGC 300
 CCNTGAN 307

SEQ ID NO:6168

SEQUENCE LENGTH:74

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07328

SEQUENCE DESCRIPTION:

GATCAGAAAA TGTGTGACCT TTTTAAAGTT AAATTGTTAA AAAATTTTAA CTATTAANCA 60
 TTTTAAATAT TAAA 74

SEQ ID NO:6169

SEQUENCE LENGTH:71

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07329

SEQUENCE DESCRIPTION:

GATCTCAGAT GCTAGAAATA GTAGACAAAG ATGTTATTTT CCAGTGCAAA TAAAGAAATA 60
 AAGATAGCAA A 71

SEQ ID NO:6170

SEQUENCE LENGTH:306

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07330

SEQUENCE DESCRIPTION:

GATCTCAATG TGCAAACTCC TTGGACTCCA GACCCCTGTC TGGTTGGCAC TCCCAGCAGT 60
 GTGTGGTTTC ATCAGGAAAT CCTTGCAGCA GGGGTTTAGA TGGTTCCTTC AGGAGGTGGG 120

NAGCCTCCAT AGCATTGAAG GCCAGCCTGC ATCTCGGGT GTGTGTCCAC TTCTCCNCTC 180
 ACTTTCCCT CACCCATCCC TTTAAAGTGA TTGGCTGTNA GGNGCACAGA GGGTGGCAAT 240
 TCCATCTGTN ATGGGNTCAG AAGGCATTNG CCGCCATTNT TGATTNACA TAAAACTGA 300
 TGNAAN 306

SEQ ID NO:6171
 SEQUENCE LENGTH:65
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07331
 SEQUENCE DESCRIPTION:
 GATCTGTCTG TTGTTGGAAT TGAGATATTA AAATCCCCTA CTATTATAGC GTTCCTCCAG 60
 TCAAA 65

SEQ ID NO:6172
 SEQUENCE LENGTH:33
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07332
 SEQUENCE DESCRIPTION:
 GATCTTCCAG CACCAAATTT AGATGGCCTC AAA 33

SEQ ID NO:6173
 SEQUENCE LENGTH:218
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07333
 SEQUENCE DESCRIPTION:
 GATCAAATGT GTGAGAAAAT CAATGGGTTT TAAAGACTAT TCTATGTCAA CTATAACATT 60
 TAATTTGGGG ATTTCTGTCC CTTATAATGT CTACCTCATT TTGGATGGAA TCCTTGAGGC 120
 CTGGTTTATT TTCCTTTNCC TTNCTACACA TCGCTGCTCA GAGTGATGAA TGGAGTTGTG 180
 TTTTGAATAA AATATCTATG CATCCNTTTG TGAGCAAA 218

SEQ ID NO:6174
 SEQUENCE LENGTH:186
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07334
 SEQUENCE DESCRIPTION:
 GATCATGTTT TAACTAACTT CATTTTGTG CTGAATCATA ATNNATGCTC TAATGACCTT 60
 TTCTTTCTTA TCCCAACCCT GTGTAATCTG CTCTATTAG GGCTAATTGT TCCTTGCTTT 120
 GTATGTATGT AATGTGTCCT NGTNTTCTTC CTCCTATTNN NTNCTCATAA AACTCCATCC 180
 NNTTTN 186

SEQ ID NO:6175

SEQUENCE LENGTH:152

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07335

SEQUENCE DESCRIPTION:

GATCCATTCA ACTTGTGTGT ATCAGTAGTT AATTNCTTTG TATTGGTCAG CATTCCATTA 60
TTTGGAATTT ATACAGTATG TTTATCCATT CACCTGTTGA AGGACATTGT TTCCAATGCT 120
TATTACTAAT AAAGCTGTGA TGAACATTCA AA 152

SEQ ID NO:6176

SEQUENCE LENGTH:236

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07336

SEQUENCE DESCRIPTION:

GATCTGTATT TAATGAGGAG GAGCCGAGGC CCCAGCTTCA TCCAGCTTCA ACCAATGCCT 60
GGACCTGTCC ACCTGANAGG CCCCTGGGGC TCCCAGCTG CTGGCCAGAC CCTGGCGCTG 120
CCACAGTCCT GGCAGTCCC AAGGCCATAC CTGNCTAGCC CTTTGGNTCC ATCCTGTGGA 180
TGCCCACTCA CCCCTNAGAC TCCTNCTGNC CATGCTGTGG NCGGACTTGT AAGCAN 236

SEQ ID NO:6177

SEQUENCE LENGTH:148

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07337

SEQUENCE DESCRIPTION:

GATCATGCTT GTGTTTACCC TCTAAGCTGA AGCTGCTCAT CAACGGTGAG ATGGCAAAAA 60
GGTGGGTCCA GAAGAGGGGA AAAGAAGGGA GTCTGTGAAA ACAAATGCT GAAGANTCTG 120
CATCAAATAA ACCCTTCCTT CCTTCAAA 148

SEQ ID NO:6178

SEQUENCE LENGTH:155

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07338

SEQUENCE DESCRIPTION:

GATCTCCTTA TTGACCACAG ATTATTCCTC TGTTTCAGCAC TCCTTATTCT TATCCCTGTT 60
ACAATTCCTA TTATGGTGCA CAAGCTCCAT AAATATTTAT AAAACTGATA TTAATTGANT 120
AAAATATTGT CTAGAGGTCT CAGTCATTCT TCAAA 155

SEQ ID NO:6179

SEQUENCE LENGTH:198

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07339

SEQUENCE DESCRIPTION:

5 GATCATCTAG ACATAATTNC TGAAGATGAA CTTTGGGACC AACTTCTAAG TCACACAGCA 60
 TCATTATCAG ATTATTACA CAACGNCTTT TINGTTTNA CTCTATTCT GAGGAAAAAG 120
 CCTTCCCGAA ATCCGTAATG ANTTTNTCCA TGGTAACCCC TCTCTGTTT TNACACAGNA 180
 AAGTTTCTCT NGA CTGTN 198

SEQ ID NO:6180

10 SEQUENCE LENGTH:188

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07340

SEQUENCE DESCRIPTION:

15 GATCGCTGGT TTCCTCTGCA GCGCGAGGGC TCCGGCGACC AGAGGATTCT NCCCGGAAGG 60
 CATTCTGCC GCGCTCCNG GGGCACCCCT CAATTGTGTA CTACGTCCTT GTTGTGTG 120
 TATCCGTGCC CACGTAGATG ATGTCTGTAA CGNAGTTTG TTTGAAATAT GAGNATATGC 180
 GGCTTAAA 188

20 SEQ ID NO:6181

SEQUENCE LENGTH:135

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

25 CLONE:HUMGS07341

SEQUENCE DESCRIPTION:

30 GATCAGTTTA TTATTGATAC ATCACTCCAT TAATGTAAAG TCATAGGTCA TTATTGCATA 60
 TCAGTAATCT CTTGGACTTT GTTAAATATT TTNCTGTGGT AATATAGNGA AGAATTAAAG 120
 CAAGAANNTC TGA AAA 135

SEQ ID NO:6182

SEQUENCE LENGTH:226

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35 CLONE:HUMGS07342

SEQUENCE DESCRIPTION:

40 GATCAGTTTT CCNAGTCTTT GGGAAACGGG TTGTGTTTT TNCCCTTGC CGTGTGGAT 60
 TTCCGGTCCC GGCTGACACC TCCCTTCGTG TGTGCGNCCC TCCCACCCC TGCCTTTGTN 120
 TCGGTGCCCT GTGTCCGTT TCTNATGTGG GGATGTTGT GCTGCAGACA AAAAGAACAA 180
 NAAAAANTTT TAAATACCA CAAAGATGGN ANAAAANGAC AAAANN 226

SEQ ID NO:6183

45 SEQUENCE LENGTH:97

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07343

SEQUENCE DESCRIPTION:

50 GATCGTCACA TGCCGGGAG CAATGTGGAT GGCCTGGGA CTCCTGGGT TTCTCCCTCC 60

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CGACTCCCTA ATAAACCCCG TGAACCTTGG AGCCAAA

97

SEQ ID NO:6184

SEQUENCE LENGTH:321

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07344

SEQUENCE DESCRIPTION:

GATCACCTTN AGTNCNTNAGC AGGGACTAGG CTTGCAGGTA ANATAATGGG CCAGGGCACC 60
 CAGTCCAGAA GGAGCAATGG CACCTGGGCA GTGCCAGGCC TTAAAGCCCG CTGCTCCTTT 120
 NCGGTAGAGG AGAGGCCCAT CACTGGTGTG GTGGGGTGGG CTCTCCCTTA GGCTTGGGCA 180
 AGGCAGCCAC CTGCCCTTNC TCTCCCTTAG TGTCCCTGG CCTCCCTGCC ATCAGGTTGC 240
 TGGGAGTGGA GATGGAGGGA TTATTGAGCA GAAAATNAGT TGGATGGAGN TAAACAGNTC 300
 CCATCCCTGG GTAATGGATT N 321

SEQ ID NO:6185

SEQUENCE LENGTH:292

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07345

SEQUENCE DESCRIPTION:

GATCAGCTTG GAAGCTTAAA GTCCAATCCT TACTTCGAGT TTGTTTTGAA AGCAAATTAT 60
 GAATATNATG TTCAGGACA AATATAACTT TTNCTAAAAA TGGCCATTGT TTATGAAATC 120
 TGTATAAGTG TGTCTTATA CAAATTTNAG GCCATAAACA AGTGTAAGTT TGTACAATTT 180
 CATAACATGT ATAGCTGAGT TTTTATACTT TATATGTAGG AAGCTAATAT AAAATAGTTA 240
 TGTAAGTGTG ATTTTGGTTT TCAGTTATGT GACTTGT TTTT TNCACCNCA AA 292

SEQ ID NO:6186

SEQUENCE LENGTH:155

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07346

SEQUENCE DESCRIPTION:

GATCCATAAC AGAGATNCAN AGAGGCACCG TGGAGTTCCA GGGTCATCGG TCAGCGAGGA 60
 ACAAGGAGGG AAAGGTGTCT TCCTGCCCTT TGATGCTCAA CTAAGCATCT GTTCCCTAGA 120
 AATACATGTG TCCAGGNNGN NTCCATGGGC TTTNN 155

SEQ ID NO:6187

SEQUENCE LENGTH:301

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07348

SEQUENCE DESCRIPTION:

GATCTTNAGT TGTTTTAATT TTTTTTTTCT GCATCTGAAT CTGCATGATT TCCAAACCC 60
 GTACCATCTG AATTTTGCAT TTNAGCACTT GCACTATTAC TCAGCAGCAG TAACATGGTA 120

ACACTTAAAA TGGTACTCGG GGACCTCCAA AGACTAAACT GNCANGCCTT CAAGGNGCCC 180
AGGGGTACGT TANCCTGTCA ACGGCATGGT TTAATCCCTT CTTTACACTT GTGTAAATTT 240
CAGTTACTGG TCATAGNNGG CTTTCAATGT TGCCTGGCCT NTTATTANCN TGTTTATGGG 300
N 301

SEQ ID NO:6188

SEQUENCE LENGTH:203

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07349

SEQUENCE DESCRIPTION:

GATCTTTATC AATATTCTGG ATTAGACAAC AAATNACCTT TCTGGGTGTT TCTTGTAAC 60
TATACTCCTG TTTGAATGTT AAACCTTGT GCTAAAGTTT AATTTTAAGA TGTTTGAATG 120
TNCAGTTTAT GTATTTGANC TACAATAAAC CAACCCCTTT NATATATCTG TNGTATAT 180
GATTATNGTN ACTTAATTNT NAN 203

SEQ ID NO:6189

SEQUENCE LENGTH:316

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07350

SEQUENCE DESCRIPTION:

GATCAAAGCA GGTGGGGCTT TGGGATGTGA AGTGGGTCTT NACGCGGGGC CCCACAGACC 60
TGTCTCCGCG TGANTGCCTG CAGGCCCTG CGTCATTCCT CAGCTGCGTG CGAGGCCTCT 120
GGCTTGGAGA GCCTCCCCGC TGTGCACAGT TCTCTCTTG CCAGGGCAGG GCTTGGGGCA 180
GCACCAAGCT GGGGCAGAGG AGGAAAGCAA GAGGACAGAC CTCCAGAAGA GCAGCGGAGG 240
GCGGGTGAGG ATTTCCCAT ATACCAGTGT GTGTTTCATCT ACCCATGAGG AATAAAACCA 300
TGCAGCAGCA TGCAAA 316

SEQ ID NO:6190

SEQUENCE LENGTH:129

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07351

SEQUENCE DESCRIPTION:

GATCCCCAGN ACTGCAGCCG AGCCCCGGGC TTCCTTTCTT ACCATTCTGT ATGCTTCCAA 60
GGTGTGACCA TTCAAATAA CNNTATTATT AAGATTATTA ATAAAGATTT CTTTCTTCAA 120
ACCAGGAAA 129

SEQ ID NO:6191

SEQUENCE LENGTH:293

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07352

SEQUENCE DESCRIPTION:

5 GATCAGAATC AGAAACAGG TAAACCTCAC TCACACATTT GGACTCATTT GAACAAAAAT 60
 CTAGGCCAAA ATACTGAAAA GCCTATGTGT TTTTAAATT GGAAGTATAT GTAAGGTTAA 120
 TGCATTTAGT GAACGTGACT AACAAAGACT AATGTGCACA TTAACAGNTG TNCTTTTAA 180
 GGTTTTATGG GAGGCTGTGC ATTGCTCAAA AGCTGTTGGG AACGCCTTCT GACCAGTTGC 240
 CTTCAGAACT AGTTTNGCT GTCANTNAA AANCCAGTGN NTTTACTCNN GGN 293

10 SEQ ID NO:6192
 SEQUENCE LENGTH:52
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07353
 SEQUENCE DESCRIPTION:
 15 GATCTGCATG TGTGACACTG ATTCTTTGGA AATAAGAGT GGAAGCTGCA AA 52

20 SEQ ID NO:6193
 SEQUENCE LENGTH:236
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07354
 SEQUENCE DESCRIPTION:
 25 GATCTAAATT TAAAAATAG GAATGAATCT NTAATTGCA GCTTGTGGAG AGTATAAAAC 60
 ATTTTGAGCA CAATATGCAG CACATAACTT TTCAGATTGT GTTGCATAG AAAAGGTTAA 120
 AATCAATTCT TTCAGTTACT GTATTGTCA TTTCTTGTA AGGTCTTCT ACATTGAGA 180
 CGAGACCCCT TAACATAGGC TGCCTTAGTA AAAAAATAA ACAGATGTGT CATAAA 236

30 SEQ ID NO:6194
 SEQUENCE LENGTH:282
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07355
 SEQUENCE DESCRIPTION:
 35 GATCTAGAGG GNGGATTCC AGCCAGGGCT GCTAGACGGA GGCCTACTCT TCCATCTTTC 60
 CTGATGGCAG GATGGCCTGG CCAGGGCCTG GAAGACAGAG ACCTCCTGCC TCCGCCTCAG 120
 TAAGACGACA AGGAAAGGCA AATGCCCAAG GGAAGAAAA GGAAGGCTCT TCTCCCAGA 180
 GTTCCCCATG CAGACATGAG TGGGTGCTCA GTTCAGANTC ACTCCGAGA ACTCATCCCT 240
 40 AATGCTGCAG ATTTGGGCTG GAACAGATTC AACTGTCTG TN 282

45 SEQ ID NO:6195
 SEQUENCE LENGTH:158
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07356
 SEQUENCE DESCRIPTION:
 50 GATCTTCATC TCTGTTTGC TCCACTTGTA TTCTTTTTC AGTAACTGGA TTAATGCTTT 60
 GAGCCCAAAA TGAGATATAT TTNTAATTAC TGTAGATTAC ATGTTGTTG GCTGGGAATA 120

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CAATACAATA AACTTTATTT TGCATAATGC ATTTCAAA

158

SEQ ID NO:6196

SEQUENCE LENGTH:113

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07357

SEQUENCE DESCRIPTION:

GATCTGGATT ACTATGTAAA TTCACAGNAG TACCGCTAAT ATAATTTTGG TTGCNTGTAT 60
 TANCATCATC TGGTCTGNAA ATGTGGGTTT TTATTGGCA CATTANATA AAN 113

SEQ ID NO:6197

SEQUENCE LENGTH:323

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07358

SEQUENCE DESCRIPTION:

GATCTGGAGC CACAGTCTGT NTATCTTCCA GTTCATCTCA GTCCTCGAGA AAGGCCCTNT 60
 AAATATGTAA CTTTCCCAT TNCCTTTAAC CATGGGTTGT GTGAGCCAGA AAGAGCTTTG 120
 AGAAAGATGG CTGCTTCCAC CAGGGNGGAG GCTTCTNGGT CTGCATGATG ATGGGGCCCG 180
 TTTCTGGCCA NAGGTGGCT CTGGGAGCAG TTGTNCTGCG GGCTTGCTGG GGGCGAACTC 240
 TAACTGTTGC AGAAACAGNG CTTNATGGCT TGCTNNGGGT ACTTAGCTGG AATATTTTAA 300
 AGTGTCAAGT AAATGTGATN TNN 323

SEQ ID NO:6198

SEQUENCE LENGTH:50

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07359

SEQUENCE DESCRIPTION:

GATCTGTATG TACTTTTAGT TGTATTAATA AAGTGACAGT GAGTGTCAAA 50

SEQ ID NO:6199

SEQUENCE LENGTH:149

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07360

SEQUENCE DESCRIPTION:

GATCACCCAG TCCTAAGGTT TCAAAAATC TTTGACATTA GATTCACAA CTGCACAATT 60
 GANCTTATTG GCCTGTAAC TATTNCTAA ATGCTCAGTG CTATTTATAT ACTACAGTAN 120
 TTNCCTGTTA AGNNGGCAGT TGTAAGNN 149

SEQ ID NO:6200

SEQUENCE LENGTH:329

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07361

SEQUENCE DESCRIPTION:

5 GATCGGACTG GAACTGTTTCG GCTGCNACCA GAAATTNATT TTCCTGAGTA AATTGCCGAG 60
 AATTAAGAAT GAAGAGGGCC ATTTGCATCT CCTTAAATTA TTCAGTTACC TGCTTTATTG 120
 CTCCATGTGG AAAACTTAAA ATTGTAAAGT TGTGCATTAC TGTATTTNAN CTTGTTGCTT 180
 AGTTTCTNCA TGTTTATTNN CAGTAATGGC TGANAGTGTT AACTGTNCCA TACTTTTNGC 240
 10 ACAATGTNCT GCATAAGGTT ACCTGTGTAC AGAGTTTTAC TTTAGNTTAN CTAATAATATT 300
 GCCTGGGTTT AGTNTTNATT TCCATTNTN 329

SEQ ID NO:6201

SEQUENCE LENGTH:150

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07362

SEQUENCE DESCRIPTION:

20 GATCTAGGAT GTGGNTGTGG AGTACTTAGC ATCGGAACTG CAATNTNAGG AGCAGGGTTG 60
 TGTGTTGGAT TTAACATAGA TGAAGACGCA TTGGAAATAT TTAATAGGAN TNCAGAAGAG 120
 TTTGANTTAA CAAATATTGN CATGGTTCAN 150

SEQ ID NO:6202

25 SEQUENCE LENGTH:142

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07363

SEQUENCE DESCRIPTION:

30 GATCCAGGCG CCTGGGCAGA GCACCTGCTG TCTGCCCTGC ACTTTGAAGG GTTGTCGTGG 60
 GGATTGAAAT NAACCAACTG CCGGCTGAGC GGAGGCTCCC TGGGAGTNAC AAGTTCCGTA 120
 TTAATAATGCT TTCAGTGGNA AA 142

35 SEQ ID NO:6203

SEQUENCE LENGTH:294

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07364

40 SEQUENCE DESCRIPTION:

GATCTATAAA ATTTGAGTTT TAAAAAATGT GTTAAGGCTT TGTTTTGGC CTAACAGGTG 60
 GTCTATAAAG AAGNAAGTTG TATGAGGTAA TGAGAAGGCT GTGTATCCTG ATGTTGTCGA 120
 GGAGTGTCT CTATACCTTC ATTAGAAATA ATTGTNTTIT TACTGCCTTC TAGTCTTCTG 180
 45 TTTCCTTACT AATATNCTCT CTTGTNTTAT TACAGAAAGT GGNGTATTGC AATATCCTAC 240
 TATAATTGTN TTGCNCTCTT TGTGTNATC CCAGTNTNNG TNAGTNTANG CTTN 294

SEQ ID NO:6204

SEQUENCE LENGTH:306

50 SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS07365

SEQUENCE DESCRIPTION:

5 GATCCCAGCC CGGCCTTTGT GGGACTAAGG TGGCGCTGAG TGGGAGAGCA GCAGGCGATG 60
 CGCGAAAGGC GGGACTTGGA GGTTCTGGAT GTTCCTGTGG ATAACCTCAA GGAGCACGGN 120
 ACCTGGGACA TGACTGCCCC CAGCATTCCT TCTGGCGGCA GACAGNGTTT GTNAGTTCGC 180
 CAACATCCTG CTTTCATGNG AACAGTTTGC TGTTTGCTNA CAGAGCCTCC AGTNGTATAC 240
 10 TGAGTTGGTC ACGACNTNA TTNCTTCGNN CCTACAAACA TNTCCACTT TTCGNTTNTN 300
 TGGTTN 306

SEQ ID NO:6205

SEQUENCE LENGTH:146

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07366

SEQUENCE DESCRIPTION:

20 GATCTTCACT AAATATTTTG AAATGAATTC ATTTATGGAA GAATCAAANC TAGGTCAGAT 60
 TGTCCAAACC AGACTAAGAA ACTGCATAGG GTNTTTTATG AGGTNTTTAA TGACTCATCA 120
 GCNCCATTAT ANCTGANATT NCTACN 146

SEQ ID NO:6206

25 SEQUENCE LENGTH:298

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07367

SEQUENCE DESCRIPTION:

30 GATCTCACTG CCCAGGCTGG TCTTGAATC CTGGCCTCAA GCACTCCTCC TGCCTCACC 60
 TCCCAAGCG CTGAGATATA GACCTGAGCC ACAGCGCCTT GCCAGTTGTT TTATGTTCTT 120
 TATCGTTTTA CCTGCTTGCC TTCTGTAACC TGGGTTGCAT TAAATTTACT CCTACTNGTG 180
 AGTAAATGTG AACCATAGCA ATAAATTTAG TCTATCCCA GAACCTTCAG TAGCAAGGAC 240
 35 CNATATTNCT ACGGTATTTG TNTGTGTCCA NANANAAATA AAATTACCTT NAAAAANN 298

SEQ ID NO:6207

SEQUENCE LENGTH:28

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07368

SEQUENCE DESCRIPTION:

GATCATTAAA GATAAACATA TTTTAAAA 28

45 SEQ ID NO:6208

SEQUENCE LENGTH:205

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

50 CLONE:HUMGS07369

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SEQUENCE DESCRIPTION:

GATCCTGTTG GCAANGGGAA TG TAGAGCTT CCAGGGTGCA TTGCCCACTG TTCTCATTGC 60
 ATCCCAGTCC TCCAAGTCAG TCTAGCCTTA CTGTACACTT GACAAGTGCT TACTCAGCAA 120
 GTCCAGACC CACGNCCTNT TATCTCCNAA GACTGGNTTT GGCCATGNCC TTCCTCATCA 180
 GGCTATGCG CANNANTATN TAGGN 205

SEQ ID NO:6209

SEQUENCE LENGTH:302

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07370

SEQUENCE DESCRIPTION:

GATCCATTTA ATATGTTATC CCCACTAATT AATTTTCGTA TATNATTTCC AATATTTGGA 60
 AAGCTCTTTA TAGCCATTG GTATTCCTA TTACCCACCT CCTATTTTAA ATATTTATCA 120
 GTCTAAACTT GTGCAGTGA GTAAACATGC AAGTTGTTAT GCTTGAGCTG TATTACCATA 180
 AGTNGCNTTT TANGTAACT GGTGNNTTTG GGCCATAGAT GTTTTGCTT TTTGTTTGAT 240
 TTTNTNTTNC AGGCTAACTG TTNGNGGTAT ACATNTATTT NTCTGTTGTA CAGATTTGAA 300
 NN 302

SEQ ID NO:6210

SEQUENCE LENGTH:301

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07371

SEQUENCE DESCRIPTION:

GATCCCCCAC ACCAGCGCCC AAGAAAGGAC CAGCGTTGCA CAGCCCGGTA GAAAGGGTTT 60
 ATTTATGCGC AACGGTTCAC ACAAGCCTTC CTGAAATTCC ACTTTACAGT AAATAAAGCT 120
 GTGCGTTTCC CTTCCCATG CACAACATGC TATCAATCTA CAACTGTCAT TTAAGTGTGA 180
 AAAAATAGNG CGTCTCCCT TTTGTCATCG TTCTGGTAAC ATTTGGCGTA GCATCTGACA 240
 GCACGGAGNT GCTCACTCCT GGACCGGTTA TTTGGTTAAA ACCCAAAATG TTAGGTCGGA 300
 N 301

SEQ ID NO:6211

SEQUENCE LENGTH:111

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07372

SEQUENCE DESCRIPTION:

GATCCATTTA ACTATTCATC AGAGGTGCTG TAATTTTAAA TTGTCTCTTG TCTCCTTCAG 60
 TTAATTTTCA GAATTAAGAA CATACCATGG GNAAAAAAA CNCNAANCAN N 111

SEQ ID NO:6212

SEQUENCE LENGTH:206

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07373

SEQUENCE DESCRIPTION:

5 GATCCTGGAA ATNAGATGGG CCTTACTCTG TCGACTAAAT GAATAGCTAT TTTCTTGTCA 60
 TTTTAAAG TGCAACTCTT GCTTCATGCT GCTTAAGTTA CCAGATGAAT GCTGNGAAAT 120
 AAGTAATCAC AGNCATTTTA ATACCATTTC ATTGCTGTTT TACGCGTGTT CATTACTTAA 180
 CAAAAAATTA TCTTTTAGCT TTAAAA 206

SEQ ID NO:6213

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SEQUENCE LENGTH:101

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07374

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SEQUENCE DESCRIPTION:

GATCTAGCCT GCCCTGCCTG CCTGCCCTGG GCGATGAGGT ACGGTGGGGA AGGTGCCTAT 60
 TTTAGAGAAC TTTGTCACAG TATTAAAGTT CCCAGAACAA A 101

SEQ ID NO:6214

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SEQUENCE LENGTH:252

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07375

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SEQUENCE DESCRIPTION:

25 GATCTAGAAT GTAACATAAT TGCTTATAAA TGTCAGAGCA TTTTNAAG GTACAGTATA 60
 TGGGGATTGT NTCGTTTTTC CTAGCAGGGG AACCTTAGTT AATANTAAAA TACTACTTAT 120
 TTGAGTIACT GNTACAGNTT CATTAAAGGC TTGTGTGCAA ATGTGGTCTC ANTCTTTTT 180
 CCCTCCATGA TTTTCCTATG TGCTTCCTCT GGCATTCCT GNGGTTTTGG TAAANTAANT 240
 30 TGCCNNTTAA NN 252

SEQ ID NO:6215

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SEQUENCE LENGTH:257

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07376

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SEQUENCE DESCRIPTION:

40 GATCAGGAGT ACCTCCCTCA AGAGTCGAGG GNGGTAGATT TCTACACGCA GAGGAGGTGC 60
 TGA CTCTCA GGGCAGGCCA GCAGTGNNT GTTGCTTTN TAAGGGAGGG CCACTAAGAA 120
 AGTTAGTGA TTGCAAAGTT GGTGCTCTGG TGGNAGTTAG ANCAGACCAA CATTTTATGC 180
 CATAACTTTA NGAGTTCCAT AGGTGGCGGC GCAGTTCTTN CGNATCAATG TNTAACTTTG 240
 NGGGGGGTGT TCATTN 257

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SEQ ID NO:6216

SEQUENCE LENGTH:189

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS07377

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SEQUENCE DESCRIPTION:

GATCAGACAT AAACCTAGCA TCTTAATGGA AGAAAAATGA GGGGAACCTC AATTATGATT 60
 TATTAAAGAC AATTTCATT ACACCTCCT TTATGACAAG TGACATTTTA GATGTAAAAG 120
 TAAAANCCTT ACCATGCCTT TTTTTTTTT NGTGGGCCTA CCTTNGGGGC NTNAANCCCN 180
 GGGGNCCN 189

SEQ ID NO:6217

SEQUENCE LENGTH:272

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07378

SEQUENCE DESCRIPTION:

GATCTCGCAA TGTCTGCCTT AATATTCAC AATTAAACCT TTCATATTTA TTATAATTG 60
 GTAATATTC TTAATTTCC TGTCTTTTA AAGTCTTGAT AGATTGTCA AAGTATATGT 120
 NCTTTCCCT ATGCTGGTTT AGAAGTNCGA NATATCACTT CTAATAGTAA TTATGCTCAA 180
 NTAGGTACAC ATATACATAC ATNTTCTCT CNCNCTTTAT ATACTTNTGT GCAATTGGCA 240
 CTTTGACGT ATGTCATANN NNTGTCATNC CN 272

SEQ ID NO:6218

SEQUENCE LENGTH:74

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07379

SEQUENCE DESCRIPTION:

GATCTCGGC CTCCTCAAGT GCTGGGATTA TAGGCATGAG CCACCGTGCC CAGNCTAATT 60
 TCTAATTAAT TAAA 74

SEQ ID NO:6219

SEQUENCE LENGTH:303

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07380

SEQUENCE DESCRIPTION:

GATCTCGTCT GCTCACCTTA GCTTTCTGCC TTGGCAGCAC GGGCTGCGGA AGAAAGCACG 60
 CTGGGCCAGG AGGCAGGGGT GCCCAAGCCA CAGGGAGCCC CTGGGAAGC CTGCTCCATT 120
 CTTCTGGTGA CCTTGGCGCT CCTTCACTCA TCTCCCTGC CCCCTCAGGA ACTGGTGGCC 180
 CAGCTTCCAC ACCNNACCT CCCAGTCTCT AGNCTATCCA TCTGTCTGTG TATGGCCTGG 240
 AGTCACTNCT TCCTCANGCC CCCAGGGNAA NGTGTGCTCA AAATAAAAC CAGNGGNTG 300
 AAA 303

SEQ ID NO:6220

SEQUENCE LENGTH:205

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07381

SEQUENCE DESCRIPTION:

5 GATCCCCTTT CCCACATCCG TGCTGTGTCA TTGTTGCTCT GCTTCCTTTC AATGTGTCAG 60
 TGCCTGGGGG GAGGGGAGGA GCACCCCTC AGCCCCCTG AACCTGACCA AAAGCCATGG 120
 CTGTTGCTCC NNNCTTTGTA TGATGCAAT GCTGAAATGT ACAAATCAA CCATGACAAC 180
 AANGAAAAAG GCCTGTACA GCAAA 205

SEQ ID NO:6221

10 SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07382

SEQUENCE DESCRIPTION:

15 GATCATAAAA ATGAAGGAAA ATGCCAAGC ATATCCCCAA CCCCTCACC CACAACCACC 60
 ACCACCACAC ATAANCAAGT TGCTGACATA NNAGGGGAC CTTTTNANN 110

SEQ ID NO:6222

20 SEQUENCE LENGTH:248

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07383

SEQUENCE DESCRIPTION:

25 GATCGCATCT CATTGTAAAC TCTTTACTGA TATGTTTGT AATACAGAAG TGAAATGTGG 60
 ACATAAAATA GTTACGCTAT TTGGTTAATG GTACTAGACA ACATGTAATT AATGACATT 120
 AAAAATTAT GGCTAGTGAT ATATATAANG TAAAATTTT TTTGCAGTAA AATATGCCCT 180
 TNATTATNGN CGGGNGGCTA TANGGNCCA ACAGTTTGT TGAAANTAGC TCCNAATCAT 240
 ATCTNTNN 248

SEQ ID NO:6223

35 SEQUENCE LENGTH:145

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07384

SEQUENCE DESCRIPTION:

40 GATCCCTTTA CACTCTTCAA AAGTGNGAAC CCCAAAGAGT TTTATGTAT GTGGGTAAAA 60
 TCTATTGATA TTTCTGTGA GAATTTAAAA ATTATGTAA AANAATAAGA NGAATGACTA 120
 TTTNCAAAA CAANANATGA GNTGN 145

SEQ ID NO:6224

45 SEQUENCE LENGTH:129

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07385

SEQUENCE DESCRIPTION:

50 GATCTGCTTG GTGAAGAGTT CAGGAAAGTA TATCAACAAA GGNGTGAANT TTGTGAAGAG 60
 AAATNNTATC ATCAAAATG GGTAAAGCAT AGNTTTTTTG TATGTTACCA CTAGATGTCA 120

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GCATANCTN

129

SEQ ID NO:6225

SEQUENCE LENGTH:212

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07386

SEQUENCE DESCRIPTION:

GATCAGAAAA CATTGATTTA TATTTTAAAA TAGGATATTG GAGCCATTGT CTTAGATAAT 60
 TTTCAAGGCA AGANCAAGCC TGCAATAAAT ANTGAGNCTT TGACATTCCT TGGCGCGGAC 120
 GATGTCTCCT GCCCTANTAT GACTGTCACT CAGCAATTC AATTTTATGN NTACTTGATT 180
 TAAANCGGGA GGCTANACTT GNTCTGAAAN GN 212

SEQ ID NO:6226

SEQUENCE LENGTH:235

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07387

SEQUENCE DESCRIPTION:

GATCTTACTG AAGATGATGT TCCAGCAGCA GCGACTTAGC CCCAGGAGCC CAGTTTCAAT 60
 GGCCTTGCTG TGTGGTGTTC CAAGTGCATT TAAAATGTGT GACACAGNAA CGGCACACTC 120
 TTNCACATGC TTTTGNAGTA TTATAANNCA CTTTATTACA AATTGTCTT NGCTATTAGC 180
 AAATANAAC TATTATCATT CTTTATTAAC CCACCTNGGC CTTTNNAAAN CCTCN 235

SEQ ID NO:6227

SEQUENCE LENGTH:290

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07388

SEQUENCE DESCRIPTION:

GATCTCACTC TTNCCCAAGC TGGAGTGCCG TGGCAAATC ACAGCTCACT GCAGCCTTGA 60
 CCTCCTAGGC TCAAGCAATC CTCCTACCTC AGCCTCCCAA GTTGCTGGGC CTGTAGGTGG 120
 ACACCANCA GGCCTGTTGA TTTTNAATTT TTTTATATAG AGTTGAGGTC TCGGTATGTT 180
 GCCAACCTAG TCTCAAATC CTNGGCTCAC GGCTGTAATC CCCAGCTCTC AGGGAGGNTA 240
 AGCGGCGGNN GGNTAGCTTA AGCCNAGGAN TNCGAAGNCC TGNCTTNGGN 290

SEQ ID NO:6228

SEQUENCE LENGTH:249

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07389

SEQUENCE DESCRIPTION:

GATCCATGGG GCCCAGGNCT AGAGGGCATC CGAGGGGCTG GGCCTTGGCC ACAGCCTCTT 60
 GGATGATGTC CTTGCTCCAC CACGCAGGCC CTTGCTTCTC CCATCAACGA GTCTTGGCCC 120
 AGGGGACACT GACCCTCCCN CAGGGAGGAG AGAGGAGTTC CTTNCCTTTA AACGAGGGCT 180

TCCACGACCT CTAAGGGGTG GCGNGGAAGG GAGGAGGATG AGAAGATGNA GATTAAATAA 240
AGGTATGTN 249

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SEQ ID NO:6229
SEQUENCE LENGTH:125
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07390

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SEQUENCE DESCRIPTION:
GATCTATCTC AGGGCTTGTT GTAAGAATGA AATGAAGTAT AAAGCCATGT AGAGCTTGTA 60
GAATATGCCT GGCACTTTGT AAGACCTTAA TANNTGTAA CTATCTAANC ACANAACAAN 120
CCNCN 125

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SEQ ID NO:6230
SEQUENCE LENGTH:71
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07391

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SEQUENCE DESCRIPTION:
GATCCTAGTG CTCCTGTCTG GTCAGGGCCT ATCTTTATGT GTTCGTAA CTTTAAACA 60
TGAGNATTAA A 71

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SEQ ID NO:6231
SEQUENCE LENGTH:182
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07392

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SEQUENCE DESCRIPTION:
GATCTGAAAA GTTATTAAGT TCAACTCCAC AGAGGTTCTT AGGAGAATGT TAACAGTCCA 60
GACCTTTTTC ACTACATCAA CCACGGATGA TTGGCATATA AAATCCTACT TAACCTTTAC 120
ATAACTGCAA TACCAGGAAA CAAATATGTC AATAAAAGGA AACATCATT CATTTTTGG 180
AA 182

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SEQ ID NO:6232
SEQUENCE LENGTH:233
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07393

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SEQUENCE DESCRIPTION:
GATCCAAAA ACCATTACAG CCAAGTNACC CTGACATGTG ACATCTACCT CCCGACCTAC 60
CACCCCACTG GCTGGTTCCA GAACGTCTCT CACCTAGACC TTGCCTCCCC TCCTCTCCTG 120
CCCAGCTCTG ACCCTGATGN TTAATAAACG CAGCGACGTG AGGGTCCTGC TTATCCCTGG 180
TNNGGNNNCA GNTCNATCCT TGCATCANTG GGGAGGACGT GATGNGTGAG GAN 233

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SEQ ID NO:6233

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SEQUENCE LENGTH:158

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07394

SEQUENCE DESCRIPTION:

GATCAGCCTG TGCCACACTT CGCTCTTGAG GAGCTCAACG GTCTGGCAGG GCAGCAGGAG 60
GCTTTCTGGG CTCTGGNCTC CACGGATGCG TGGGCAGNGG AATGTGGGCT ATGTAGTCAT 120
AATAANTTAG GNCACAGAAA AAAAAAANAA AAAANN CN 158

SEQ ID NO:6234

SEQUENCE LENGTH:51

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07395

SEQUENCE DESCRIPTION:

GATCTCTTTT CATTGTCAA CCTTTTCAGT AAAGCCCTCT GTTACATCAA A 51

SEQ ID NO:6235

SEQUENCE LENGTH:280

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07396

SEQUENCE DESCRIPTION:

GATCAGCGAG AAGGACAAGA AGAANNTTCT GGACAAGTGT CAAGAGGTCA TCTCGTGGCT 60
GGACGCCAAC ACCTTGGCCG NGAAGGACGA GTTTACGCAC AAGAGGAAGG AGCTGGAGCA 120
GGTGTGTAAC CCCATCATCA GCGGACTGTA CCAGGGTGCC GNTGGTCCCG GGCCTGGCGG 180
CTTCGGGGCT CAGGGTCCCA AGGGNGGGTC TGGGTCAGNC CCTNCCATTG AAGGAGGTGG 240
NTTAGGGGNC TTTGTTCTTT TNGTATGTTT GTCTTTGN GN 280

SEQ ID NO:6236

SEQUENCE LENGTH:199

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07397

SEQUENCE DESCRIPTION:

GATCTAAATA CTGTGTTAAA ATATAGAAAA ACGGAGACTC TNATAAGGTT TTTNACCTCT 60
ACACITGAAA GTGAGCTCTT ACAAGAAACT AATCAGTGTT TCGTTATTCC ATATCAGTGG 120
CTTTTACTGT CAACGNTTGT GTAAGCTNAC TTATCTGGCA AACTTTT NCT TTCCAGAANT 180
ACTATCTATN TTACTNTNN 199

SEQ ID NO:6237

SEQUENCE LENGTH:279

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07398

SEQUENCE DESCRIPTION:

GATCCACCCA CCTCGGCCTC CCAGAGTGCT GGAATTACAG NCGTGAGCAC CGCGCCTGGC 60
 CAAGNAAAAA TTATTAGAAC AAGTGAGGCA TCACACTCCT CTGCCATTCA GTCAAAAGTA 120
 NGCACCAATT TATTAACACA AACATTACAT AATACATGAT ACAGGCGCAA TCTCGCTACT 180
 GCTAGCACTA CGNNAATGTT GTTCTTGAC TATAACATTA AACTGTAAAA TACTGACAGT 240
 GAAATGCACC NATANNGGNN ACCTGTCATC ACNGGGNNN 279

SEQ ID NO:6238

SEQUENCE LENGTH:21

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07399

SEQUENCE DESCRIPTION:

GATCCAAAAA CTAACCTCAA A

21

SEQ ID NO:6239

SEQUENCE LENGTH:174

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07400

SEQUENCE DESCRIPTION:

GATCCAAGTA CTCTCTCTCC TAAAGACACC ACCTTCCTGC CAGCTGTTTG CCCTTAGGCC 60
 AGTACACAGA ATTAAGTGG GGGAGATGGC AGACGCTTTC TGGGACCTGC CCAAGATATG 120
 TATTCTCTGA CACTCTTATT TGGTCATAAA ACAATAAATG GTGTCAATTT CAAA. 174

SEQ ID NO:6240

SEQUENCE LENGTH:294

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07401

SEQUENCE DESCRIPTION:

GATCTTGAAG AGACCGCTGG CAGCACCAGT ATTCCAAGA GGAAGAAGTC TACACCCAAG 60
 GAGGAAACAG TTAATGACCC TGAGGAAACA GGCCACAGAN GTGGCTCCAA GAAAAAGAGG 120
 AAATTCTCCA AAGAGGAGCC GGTGAGCAGT GGGCCTGAAG CGGCGGTTGG CAAGAGCANC 180
 TCCAAGAAGA TGCAANAGTT CCATAAAGCA TCCCNAGGAN GNTTAAGAAT GCAAATGGAC 240
 CATTCTCTTG GNNGGTTNGG NCATACCCAT NAGCCCNAGG TGACATTTCC CNTN 294

SEQ ID NO:6241

SEQUENCE LENGTH:295

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07402

SEQUENCE DESCRIPTION:

GATCTCGCCC TTACCCCGGG GTCTGGTGTA TGCTGTGCTT TCCTCAGCAG TATGGCTCTG 60
 ACATCTCTTA GATGTCCCAA CTTGAGCTGT TGGGAGATGG TGATATTTTC AACCCACTT 120

CCTAAACATC TGTCTGGGGT TCCTTTAGTC TTGAATGTCT TATGCTCAAT TATTGGTGT 180
 TGAGCCTCTC TTCCACAAGC GCTCCTCCAT GTTTGGNTAG CAGTTGAAGN GGTGTGTGGG 240
 TGGGCTNITG GGNGTGAGGA TGGAGTNTA AGTGCCNACT NNCNNCTNNN ACCNN 295

5

SEQ ID NO:6242
 SEQUENCE LENGTH:201
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

10

CLONE:HUMGS07403

SEQUENCE DESCRIPTION:

15

GATCTAATCC CTTCAAGGAA GTGATAACAC TGGGAGTGGT AACAAGAGGA GCAGGGAAGC 60
 AAGGGCCAGC CCCTGGTTCT CCATCCCNAT GTGTNGAGG GTGGGACCAG NTGGACCTGG 120
 TCCAGGCCAG GCCTCATTCC TCCTGGNCCN GGCAGGGTCA GAGGGNGGAG GGGCNTGGTC 180
 CAGGNATGGG TCCNTTCAN N 201

SEQ ID NO:6243
 SEQUENCE LENGTH:26
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

20

CLONE:HUMGS07404

SEQUENCE DESCRIPTION:

25

GATCTTGGCA GTTGATAAAA CGTAAA 26

SEQ ID NO:6244
 SEQUENCE LENGTH:324
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

30

CLONE:HUMGS07405

SEQUENCE DESCRIPTION:

35

GATCGNTGCC TGGTTTGTCT TCTCTGTGTC CTGGAGCAAA GCCAGTTCCT AAAACTAAAA 60
 CTCCATTCTA GTCTTGGGAA GAAAAGTTT TACTCAGAAC TGGGGAAGGA GTGGAACCTA 120
 TGACTTGGGC CTCTAGGCTG TCTCTGTCCC CTCAGCTCCC CGACATGCAT TTA CTCTCTG 180
 CCGTGGGTCT GCAGTCGCTG CAACCTACCC TCTCTCTGCC TCAGCCTTAC ACCCAAGCAG 240
 TAGGTCTGTG CTCTCCCTGT CTCTAGGTCG NTGAGAGAGG TGCTTTTNTT CATAAAACCT 300
 TTGGGGTTTG GGATTTCCCC AGAN 324

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SEQ ID NO:6245
 SEQUENCE LENGTH:256
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

45

CLONE:HUMGS07406

SEQUENCE DESCRIPTION:

50

GATCCATTCC GGCGCCTTCC TCTCCCAGTC ACCCAGAGGG CCCCAACCCC GCGGNCCTT 60
 TCTTCTCAA ATGTCCTCGG CTCTATACCG TGNCTGGGTC TTTTNTCTNT CTCTCTGCCT 120
 GGAAGATTCC TTCTTTNCCC TTTTGTCTTG CCCACTACTG TTTACCCCTC AAGTTTCAAG 180

55

GTCATGTCAC TGTCTNAGAG AGGTTTTCTT GTGNTNGACC TGTTTTTNAT CAGGAAGGCT 240
TGCTNNTTTC CATNAN 256

5 SEQ ID NO:6246
SEQUENCE LENGTH:37
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS07407
SEQUENCE DESCRIPTION:
GATCATTAAA CCTGGCTTGA GTCTCTGTTC TGGCAAA 37

15 SEQ ID NO:6247
SEQUENCE LENGTH:152
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07408
20 SEQUENCE DESCRIPTION:
GATCAAGNTT GNTCTTCTGT GATATATGGG ATATNATGTT TTAAAGACCA TCTTGAATA 60
CAATTAGAGA ACTTAGTATT TTGATGTACT AAGACCTATT TTAAGTTTAA TATTCTACTT 120
TGCAAAANCT TTAATTAANG ATGTTATTTA AA 152

25 SEQ ID NO:6248
SEQUENCE LENGTH:213
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07409
30 SEQUENCE DESCRIPTION:
GATCCTGCAT TAAGAAATTC AAAAATCAGT TAATCATCGT TACATGTTGT AAAAGTTGGA 60
TACTGACAGT CTCTAATAGT TTATCATCCA ATNGCANGTA ACCTACAATT TTAGGCATAA 120
ACGTTGATGT GTATAANCAAT GNATACAACG CTANTGGCTG GTCACAGTGT TTTCNCAAAC 180
35 TACAGNTCAC ACCANGTNTT TCTCCTNTTC CTN 213

SEQ ID NO:6249
SEQUENCE LENGTH:274
SEQUENCE TYPE:nucleic acid
40 TOPOLOGY:linear
CLONE:HUMGS07410
SEQUENCE DESCRIPTION:
GATCTCCTCT TAAGCGGAAA AGAGATTGAC CTTCTAGCAG AGGCAAAGAA TGGAGTCTTC 60
45 TGGGGGCAGC TGCAAAGCGT TCTCCCTAGG ACAGATGGAG CCTCCCTTTC CTCATCTACT 120
CTGTGGGTGG TTTCAGGGCC CACGAGTCAA CATNAGGAGT TGTGCTGGTG GTATGTGTGT 180
TGGAGGCTGG GCTGGCTGAT TCACAGTGAC GAGGATGTCA CTAATANCAN GTATTGCGAN 240
TGNTGATACC TAATAANAGN CTTTTTTTCC CAAA 274

50 SEQ ID NO:6250

55

SEQUENCE LENGTH:216

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07411

SEQUENCE DESCRIPTION:

GATCACCATG CTTGAATGTC CTCTACTGCT GATTCTAACA TCCGTGTCAG TTTTGGACCG 60
 GTTTCAGTAG ATGGATTTTN CTCNNNGNA TGGGCCATAC TTTCTGTCAT TCTTTCATG 120
 CCTCATAATC TTTGATTGGA TGCCAGACAT TGTAATTTT ACCTTGTTGG GTATTGGATA 180
 TTTTGTATT CCTATAAATA TNCTTGNCT TTGAAA 216

SEQ ID NO:6251

SEQUENCE LENGTH:319

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07412

SEQUENCE DESCRIPTION:

GATCTAATTT TGTATGTGGG AATCTTGCCT TGCCCTGATT TCCAGACCCA TAGGCAGGCT 60
 TGCAGGCTGT AGCTGGGGAA GACCCACGC TTATNCTGGT GCCCTGAGCA AGCCCTCCTG 120
 CTCCACTATT AGGCAGAAGC ACCTGTTCTG CCACAGGAGG CGCTGTCTTT TGCTGGCAGA 180
 GCTCAGTTTC CCTTAAGGGG AAGTTCCTTA GGATTAAAGG CCTCATGACT ATCTCAAAAC 240
 CCATGTTTGT TTGCCAATAA AGNAAATAAA TAAACTGTAA AAAANNAANN AACNACCANN 300
 NAAAAAACAC NAACANCAN 319

SEQ ID NO:6252

SEQUENCE LENGTH:111

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07413

SEQUENCE DESCRIPTION:

GATCTACTGT TTGCATGTAA GAGACCAGGN TATGTAATC TTATATTTTA AGTGTATACA 60
 TATTGTGTAT ATAACATATG GATATTAATA ATGGGGANTT GCACATTTAA A 111

SEQ ID NO:6253

SEQUENCE LENGTH:108

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07414

SEQUENCE DESCRIPTION:

GATCCCAGGA ACTGTGGGCA CCCATTTTNT GTGTCTCCA GCCNATTTCC ACTCCTAGTT 60
 TGTCATGGNT ANTTTTTGT CTTCCTGTG TGAGGGGNGC CATCAACN 108

SEQ ID NO:6254

SEQUENCE LENGTH:317

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07415

SEQUENCE DESCRIPTION:

5 GATCACAACC ATCAGCATCA CGGCCACAGC AGCCAGTGTT GGGGCTGCTG GCATCCCCCA 60
 GGC GGCTCTG GTCACCATGG TCATTGTGCT TACGTCGGTC GGCTTGCCCA CGGAAGACAT 120
 CACGCTCATC ATTGCCGTGG ACTGGTTCCT TGACCGGCTT CGCACAATGA CCAACGTACT 180
 GGGGGACTCA ATTGGNGCGG CCGTCATNGA GCACTTGTCT CAGCGGGNGT GGAAGCTTCA 240
 GGAAGCTGAG CTTACCCTAC ACAGNCTGGG GAAACCCNTA CAAGTCCCTT AATGGNACAG 300
 10 GACGCAGGGG AGCATAN 317

SEQ ID NO:6255

SEQUENCE LENGTH:218

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS07416

SEQUENCE DESCRIPTION:

20 GATCGAGACC ATCCTGGCTA ACACGGTGCT GCGGAGATA AACTCCCTT GAAGCAGTGG 60
 AGTACAATCA AACATCTTGG CTGCTCCTGA AACCGGTCC CACGNATTTC AGTCCGGCTA 120
 AGANCGAAGA GCTGGTAAAC CANTAANTTG GGTGGAGGCC GAGAGCTCGG GGCCGCGAGC 180
 AAGCCTCCGA CGANNTGGTC CCCTGGACCN NCNGGTTN 218

SEQ ID NO:6256

25 SEQUENCE LENGTH:194

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07417

SEQUENCE DESCRIPTION:

30 GATCTNACTC ACAGAGGGAA GAACCTATAA TTCTTCACAG GTCACATAGA AGCATGANAA 60
 TTTGGGTCA AGCAAGTAAA TTCTAAATCA GAATCCATAC ATAAAGTGTT TGCAATGTCC 120
 AGTTATATCT CCATGATATT TNCTTTGTGG AAGTTGATTG TCCTTCCTTA CAATAAATTG 180
 CTTGANITGT CAAA 194
 35

SEQ ID NO:6257

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS07418

SEQUENCE DESCRIPTION:

45 GATCCTGTGT ATTTCCCTGA GTCTTCTAAC ATATGAAAAT TCATATCTAA ATCAACAAGT 60
 GACTGTAATC TGGTACTATA AATACTAAAT AAACACTTCT TCATAACACT GTACCAATTC 120
 AGCTTTTAAA TTTNATTACT TTGCTTTCCT GTCCTTTGCC AACTCTTAAC CTAGTTAATC 180
 CTAGTNCCTG TGACATTGGA CCAGGCTCAG TAAATAANCG AATGGATTTC CAGCCTTTT 240
 TTCCCATCTG TNCCTGCTTT TANGTCCTCT GAATCTGCTT CTTTGCTTNC TGCTGCTTTA 300
 ATTTTANCAG TGNTTTTNGT CAAAACATAG GNTNCAGGGN CTNNW 345

50 SEQ ID NO:6258

55

SEQUENCE LENGTH:167

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07419

SEQUENCE DESCRIPTION:

GATCTGGGTT CTAGAGGGCA TGTGATGACT GTAAATGTTC ACTGGGTGGG TAGGGAGTGG 60
TATCCAGTGT TCAAGTGCAG AAATCTTTGG CTTNGCTACC AGTTCCATAT GATGAGAAAT 120
AAACGTTTCG TGAGGTTTGG TTTCATAAAA AGANNNNNAA AANNTCN 167

SEQ ID NO:6259

SEQUENCE LENGTH:130

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07420

SEQUENCE DESCRIPTION:

GATCCACACT TTAAAGAAA TTCCATAAAT GTATATTTTG TATTATGTAT TATTTCTG 60
TCCNAAGAAA ATATGTGAAT TCAGTTCTAA CTTNNGGANT GTACTGTTTG TTTTCANGTT 120
CATTGNAAN 130

SEQ ID NO:6260

SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07421

SEQUENCE DESCRIPTION:

GATCCGAGAT GGCACCATG TACTCCAGCC TGGGCAACAA GAGCGAACT CCGTCTCCAG 60
ACAAACAAAC AAANCAANAA CAAGTGTCCT TATNNGNATG GGTGTGTTN 110

SEQ ID NO:6261

SEQUENCE LENGTH:310

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07422

SEQUENCE DESCRIPTION:

GATCTGGCTG TCTAGAGGGA TTCTACGACT GCGTGCTGCC GCCTCCCCAA GAGGCATTCA 60
GGTTATTGGA GAACTAATCT CATCTCAAGG GGCCAGNCAC CAAGTCCCAA AGCCTACAGA 120
CCTCTTTCCG CCAGGCCCTG AAACCTGGCC CCGTGCCAGC AGGATGACAA GCCCCAGGGC 180
GNTCCTGATG AATATGGCTT GGAGATGNTG TACAANTTTT TATTNCCCTN TGGNTTTTGA 240
GGNATGAAAT GATTTGCACT TTTGAAAACC TGTTAACCGT AGGCCNCTTN GGNCANTGAN 300
GNNTGGAAGN 310

SEQ ID NO:6262

SEQUENCE LENGTH:129

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07423

SEQUENCE DESCRIPTION:

5 GATCCACTCG GNCAACCCAA CAGTTTTACC AACCACCCCG GGCTCGGAAC TAATAGGNAA 60
 AGTAGACTCT TTGTGAAGAA ACGAGCCANT NNCTGGAACA CCCTGGTGA AACCTGTTGA 120
 CANGCCCTN 129

SEQ ID NO:6263

10 SEQUENCE LENGTH:157

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07424

SEQUENCE DESCRIPTION:

15 GATCACTTGA ATGTTAGGNT TCATAGNACT ATACTAATCT TCTCACAAA GGTCTATAAA 60
 ATACAGTCGT TGAaaaaaAT TTTGTATCAA ANTGGGGNGG CAAATTNGAA GCTTCTCCTT 120
 AANCTGTATT GATACTGACT TGAATTCCTT TCTANAN 157

SEQ ID NO:6264

20 SEQUENCE LENGTH:171

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07425

SEQUENCE DESCRIPTION:

25 GATCTCCTTC TTAGCTCAA GGTGCTGCA ANTTGGAGCT GAGCTGTCAG TGAACGNGT 60
 CCTGGAAATN ATTAAGCAAG GCGTCGTTGC GCTGCCCAAN GACAGACTGA AGAAATTTCC 120
 AGANTTGAAG TTCAAATATG TGAAGNGGA GCAGCCCGNG GNGTTTTAT N 171

30 SEQ ID NO:6265

SEQUENCE LENGTH:174

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07426

SEQUENCE DESCRIPTION:

35 GATCCTACAT TTACAGACAG TTTTGTGATA ACCCTTTGCA TTGCAGCACC TAGTACAATT 60
 CCTTGGAAAC AGCGTGGCTC AATAAATTTT TATTGAATGA ATAAATGTGG GCCCAGAAGC 120
 GTGCTAGANG AGTGCCCTTC TGGGCTACTN AAAAAANNA AAANACNAAN AANN 174

40 SEQ ID NO:6266

SEQUENCE LENGTH:123

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07427

SEQUENCE DESCRIPTION:

45 GATCAAAGAC TCCCTGGAGC GAGAAAACAG TTAAGTCCAG AAGCAGAATG GAAGAGCCAA 60
 AAAGTACACA AAATGGACGC CATAANTTCT GAAATAAAAG TGTATGATGT GTTCTGAGTC 120
 50 AAA 123

55

SEQ ID NO:6267

SEQUENCE LENGTH:174

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07429

SEQUENCE DESCRIPTION:

GATCTGGGCC TCAAAATACC ATAGTAGCTG CTTGATAAAA TTCTAAAAAT ATCTGGTTCT 60
CTATTATGTA AACACTATTA CAGTCACCAG TGTGTGAAGA CTCTTGAGTC TGGTTCTCAT 120
ATCAGAGTCA TCATTTTCT TCCNGTGGAA TAAANTGCCT TGTGGNCTTC CAAA 174

SEQ ID NO:6268

SEQUENCE LENGTH:163

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07430

SEQUENCE DESCRIPTION:

GATCTTTTCA CTGGCTGACT CTCCCATATC TGCAAGAGAT TCTGCAGGAA CTGGGTGTGC 60
ACACGGTGTT GTAGCCAGTT CAGGTACTGN ATATTAGGA TTGGTGNCG TTCACTGTAT 120
TGCTATATTT TTGTAGCTAA ATAAANTCA ATAANTTGT AAA 163

SEQ ID NO:6269

SEQUENCE LENGTH:192

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07431

SEQUENCE DESCRIPTION:

GATCTCCAGT ACTTTGTACA ATGTATAGTA TATTAAATAC ATTTATGTGT AATTTGTGT 60
GTATTNTAT ATGGTATAAT TAAAATNATT ACATTATGT GATTGTAATT CACAAAANGT 120
ACTACTTACT ATATNCTGTA GNGTGTGTTG TGAATGTNTT TACCNGTGTG TGAGCATTAT 180
GGNGTAGGGN TN 192

SEQ ID NO:6270

SEQUENCE LENGTH:85

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07432

SEQUENCE DESCRIPTION:

GATCTTCTTT TGTAATGTTT TTCATGTTAC TGCCTAGGGC GGTGCTGAGC ACACAGCAAG 60
TTTAATAAAC TTGACTGAAT TCAAA 85

SEQ ID NO:6271

SEQUENCE LENGTH:284

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07433

SEQUENCE DESCRIPTION:

5 GATCCTAGTG AGGGATTGAT GAATGTTCTA AAGAAAATTT ATGAAGATGG AGACGATGAT 60
 ATGAAGCGAA CCATTAATAA AGCCTGGGTG GAATCAAGAG AGAAGCAAGC CAAAGGAGAC 120
 ACGGAATTTT GAGACTTTAA AGTCGTTTGG GGAACGTGTA TGTGATGTGG AAATACTGAT 180
 GTTTCAGTA AGGGAATATT GGTGAGCTGC ATATATAAAT TTGACAGATA GCTATTTACA 240
 TAGCCTTCTA AGTAAAGGCA ATGAATTCTC CATTTCCTAC TGGN 284

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SEQ ID NO:6272

SEQUENCE LENGTH:211

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

15

CLONE:HUMGS07434

SEQUENCE DESCRIPTION:

20 GATCAGTATT AACATCCAGG TTTTGATTGT TGTANTATGG TTACCTGGTA GATTACCTTT 60
 GTATATAGGN AATAGACACT GGAATAGTAA GGNGTAGTGG GTCAAGATGT CTGCAACTTA 120
 TTGTTCAAGT TCACTCAGAA NTGGGNAATC TGATATAAGG NCAAAANTGGG AGCACTGTGG 180
 CCTNTTTTNN CAACTTTGT GTACATCTGN N 211

25 SEQ ID NO:6273

SEQUENCE LENGTH:170

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07435

SEQUENCE DESCRIPTION:

30 GATCTCATGA AGTCTTCCTT GGCCCAACCA GGCAGATGCC CCGGGCTCCA GTGGGGGGAG 60
 GGGTCTGGGG TCTGGTCCGG GTTCCCNAAAT TTCTCTTNN CCCANATCNC TATTTCTACG 120
 CAGTGCGNAT AAAAATTGGA NTCGACTAAA ACCCTTGGTG CCCNGAAAAA 170

35 SEQ ID NO:6274

SEQUENCE LENGTH:301

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07436

40 SEQUENCE DESCRIPTION:

GATCTGCCCCG CCTCGGCCTC CCAAAGTGCT GGGATTACAG ACGTGTGCCA CTGNGCTCGG 60
 CCTCAGATTC CATATTNAA CACCAGCTGA TTGAGAGAAG GGGAAATNAGA AGAGCTGGAT 120
 GAGTTTAAAT AACTTTNATT GTTCAGATTC CTGAACAGGN GTTGGGATAA TGGCCATCTT 180
 TNCTTTCCTA TCCTTCCCCC CCTCACTGTG AAAAATAACA GTCCACCCCA AGTCATACAC 240
 45 TGGACCCAGT GCCTGCGGGG ACAGGNCTGT GGGTTTCTTG GTCACACCTG TGTGGTGNT 300
 N 301

50 SEQ ID NO:6275

SEQUENCE LENGTH:198

SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS07437

SEQUENCE DESCRIPTION:

GATCCAAAA ACCATTTCNGG CCAAGNAACC CTGAAANGNA ACATCTACCT CCCGACCTAC 60
 CACCCCACTG GCTGGTTCCA GAACGTCTCT NACCTAGACC TTGCCTCCCC TCCTCTCCTG 120
 CCCAGNTCTG ACCCTGATGC TTAATAAACG NAGCGACGTG AGGGTCCTGA TTCTCCCTGG 180
 TTTTNGGGGG GNTCCATN 198

SEQ ID NO:6276

SEQUENCE LENGTH:102

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07438

SEQUENCE DESCRIPTION:

GATCTCTACT CAAAGATGTC TTGCAGTTTA TGAAGTGA TGTAAGTTCA TGACTAACAT 60
 GATGACATCG TTATTAATA CATAAACTTT ATAAAGCTTA AA 102

SEQ ID NO:6277

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07439

SEQUENCE DESCRIPTION:

GATCTTAATA CCTAATTTCA TCATTTCTGA AAATGTAAGT TTTTAGAATG TATTACAATA 60
 TCAATGTGAA TATCTTGAAT CCTGTTACAA ATCCTGCACT GTATTAAACA TGTAATTTAA 120
 TTGTTTGTCT GATTAGCCAA TCTCACCACC CAAATGGGGA GGTATACATG TTTGANGAAC 180
 TGTGTAACTC AGTAATTGAT TTGTNCTGAT GTTGTAAGTC ANTAGANGTG TTTTGGAAGG 240
 NAGCATGGTG TGTGAGGCAG TGTCTGTCTT TTTGTGCCAG CTCTGTATGN TGTTTGTNGG 300
 CCNTGTTTGT AGGCCTGN 318

SEQ ID NO:6278

SEQUENCE LENGTH:321

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07440

SEQUENCE DESCRIPTION:

GATCTCGCCA ACAGCTAGCT GCTTAGGAGT ACCCCCACGA TACGCACAGC ACACCACTGT 60
 CCCTTCACTG CACTTCTTTC CTGCCTTAGG TAGTTGGGCT TGCCACCCT AGTTTGCTTT 120
 TGTAGTGGTT TGGCAAGGTT AGAAGGCCTC GGCCTCTCTG TNATGCTGGG AAGTGCCTAC 180
 TCTCTGGGCC ACTGCTGCAG AGGCCGTGGC ACTTGTNATG GGTTTGGAAG ACCCAGCCAT 240
 CTGCAGCAGA GGCAGCCTAT TCCCATTTCA AGGNGNGGAN TGAACGGAGT AATTATTCTA 300
 CTNTTCTTTT TACATAAATA N 321

SEQ ID NO:6279

SEQUENCE LENGTH:290

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07441

SEQUENCE DESCRIPTION:

GATCCTGCAC GTCCGCAAGC CGGAGAAATA CCAGCTGCAC CTCCACGAGA GCTGCGTGCT 60
 GTCCCTGAAG TTTGCCTCCT GCGGACGGTG GTTTGTAAGC ACCGGGAAGG ACAACCTGCT 120
 CAACGCGCTGG AGGACGCCGT ACGGGGCCAG CATTTCAG TCCAAGGAGT CGTCCTCAGT 180
 CCTGAGTTGT GACATCTNCA GAAATAACAA ATACATTGTN ACAGGCTNGG GGGCCAAGNA 240
 GGGCACCGTG TATGNGNTNG TNTNCTTNGG CCATTNCCC NTTCTGTNN 290

SEQ ID NO:6280

SEQUENCE LENGTH:130

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07442

SEQUENCE DESCRIPTION:

GATCCAGTCT NTGGTATCAT GCTAGTNCTG ACAGGGCCTT GATAGTCTAG AGTTGGAAAA 60
 GATGGTAAGC TTTTGTNAGN GTTTTAACAT TTNCTTGATG AAACAATAAA AAGAGGTAAG 120
 CTTTTNCCTN 130

SEQ ID NO:6281

SEQUENCE LENGTH:192

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07443

SEQUENCE DESCRIPTION:

GATCCAGTTT TGTTTGGGAC CAAAAACCAG TATTGTACAA AGTATTAAGC ATATATTTN 60
 ATATTACTA AAATNGACTG TGGTGACTTT GGATAATAAG NAAAANTTTA ATATTAAANC 120
 CATGTTTATT ACAGTATAAT TAACATGTTA AACCATGGGG NTAAATGCCA TCANTAAAAA 180
 ATTATGNCAT NN 192

SEQ ID NO:6282

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07444

SEQUENCE DESCRIPTION:

GATCCACCCC CTGCCGCAA AACATTGCTC CTAACCTCCAC CGNCTGTCCC AAAACCTATG 60
 NGANCTAATG ATAATCCCAC CACACTTTAC TGACTNTNTT TCCAGACTCA GCCCGGCTGN 120
 ACCAN 125

SEQ ID NO:6283

SEQUENCE LENGTH:301

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS07445

SEQUENCE DESCRIPTION:

5 GATCTGAGTT GCAGCTCTGC ACCCTAAATC ACACCCTGGG CATTGTCTGG GCTGCAGGGC 60
 TGCCAGGTTC TGTACTTGTG TCCAGCTATG GCCCTGGNTG CTGGAGCTGG AGGGTTTTCT 120
 GTGCTCAGAC TGTAGCCTGT AGCTCTTGGC CTGTGTAGAG CCCCTCCTG TGCCCTCAGT 180
 GGCTGTCGTT TGTTAACATN ATCAGGAAGA TGGGAAAGGT CAGGCAGNAT TTTTCTGNCC 240
 TACAAAGGGN GGNNGAGAAA GGACACAGTA TTTTNAATGA NTTTACCATA TATCTTTGGN 300
 10 N 301

SEQ ID NO:6284

SEQUENCE LENGTH:108

SEQUENCE TYPE:nucleic acid

15

TOPOLOGY:linear

CLONE:HUMGS07446

SEQUENCE DESCRIPTION:

20 GATCCTTGNT TTCTAAAAGC CAGTCCTNCC CTNCAGCTTT CCACAGTTTC TGTAATGTT 60
 TAATACTTNT ACAGTCAATG GCAATTTTAA ATATATATAT ATAANATN 108

SEQ ID NO:6285

SEQUENCE LENGTH:159

SEQUENCE TYPE:nucleic acid

25

TOPOLOGY:linear

CLONE:HUMGS07447

SEQUENCE DESCRIPTION:

30 GATCTGGGGG AAAGGCTGCC AGGGCCAAGA GGCTCGTCTC ACTCCCTGCC CTGCCATTTC 60
 CCGGAGTGGG AAGACCTTGA GCAAGCAGCA CTGCCTTCCT GAGCCCCNGT TTTCTCATCT 120
 GTAAAGTGGG GGTAAATAAC AGTGATATAG GNGTGCAAA 159

SEQ ID NO:6286

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

35

TOPOLOGY:linear

CLONE:HUMGS07448

SEQUENCE DESCRIPTION:

40 GATCACTTGA CTCTTGTAAG CGGACCTATA CTCAAAAGGC TTTGGCACAC TGTCACGNTA 60
 TTAGTATACA AATGTGCATT TGTGTTTAAG TTAAANTGAG GTNTGCNGGG TAAACTGCNA 120
 ANTAN 125

SEQ ID NO:6287

SEQUENCE LENGTH:63

SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS07449

SEQUENCE DESCRIPTION:

50 GATCTACAGT AATTTATTCT NTCTAAAAGT AAAGCATTTT CAAAACCTCAG TATTTAAACC 60

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AAA

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5 SEQ ID NO:6288
 SEQUENCE LENGTH:104
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07450
 10 SEQUENCE DESCRIPTION:
 GATCTTTTCC ACTGGNCTAC TACCTGACCA TAAACATGTC TGCATTTGAA TTCTCTAAAC 60
 CCTAAATCTG TGTCTATGAA AAATACAAAT GACTATTAAA TAAA 104

15 SEQ ID NO:6289
 SEQUENCE LENGTH:123
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07451
 20 SEQUENCE DESCRIPTION:
 GATCTACCCA CTCGGCCTC CCAATGTGCT AGGNTTACAG GCATGAGCCA TGGCGNCCGG 60
 NCAAGTTTCT ATTCTTAAAA TAAAGAGGTT ATACATCTAA NTTACAANTC GTGGCCAAAA 120
 NGN 123

25 SEQ ID NO:6290
 SEQUENCE LENGTH:296
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 30 CLONE:HUMGS07452
 SEQUENCE DESCRIPTION:
 GATCTGCCCC NCTCAACCTC CCAAAGTGCT GGGATTATAG GCGTGAGCAC CGCCCCCAGN 60
 TGATTCCAGG CGAATTCTTC TCTGATGGGC GGGCGAGGGT GTGTTCTGTG GATGGGTTGG 120
 NGTGTGTGTG TCTNNGTACA CAGATGATGT GTTTCCCTT CAGCTTCTTA CGTTTTCTGA 180
 35 GCATCCGTTG TGCCTTAACA TTTTCTGNTT GTCCTTTGGG NCAAGCAGT ATTTTACTNA 240
 TNCTTTGAAT GTTCTCATT CATTNTATCA TGTGACTTAT TAAANNAGN TTNTAN 296

40 SEQ ID NO:6291
 SEQUENCE LENGTH:228
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07453
 45 SEQUENCE DESCRIPTION:
 GATCTGATGG ATTAATAAGG GGTNTCCCN ATTGCTTGGC TCTCATTCTC TCTTGTCTGC 60
 TGCCATGTAA GATGTGCCTT TTGCCTTCCA CCATGATTGT NAGGCCTCCC AGCCACATGG 120
 AACTGTGAGN TCATTAAACC TCTTTTCTT CATAAATTAC CCAGTCTCGG GTATGTCTTT 180
 ATCAGCAGTG TGCAAGNNGG GGNNTACAC CACTNCTCTA ACTANNTN 228

50 SEQ ID NO:6292

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SEQUENCE LENGTH:176

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07454

SEQUENCE DESCRIPTION:

GATCCTGTGG TCAGAGGCCT CTTTGGCAGT GTNTTCTTAC CCTTAAGNAA GGTCATGAAA 60
TCCAGAAGGG GCAACCTTTN CAGGAGAGCT TTGGAGTCAT TTCTGTGTGA GNCACTATTG 120
CATAATCCTG NCAGNTTGCT TTTATATTTA AGGANTGATG TTACTIONCA ACTGAN 176

SEQ ID NO:6293

SEQUENCE LENGTH:310

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07455

SEQUENCE DESCRIPTION:

GATCTGACTG CTGGCTATAG GATTTTGGAC TTAATGACTG AAATTGCAAA TTGTCCTTTT 60
TCTTGGCATT ACAGATTTTG CCAAAATAAC TTTTGTATC AAATATTGAT GTGTGAAAGT 120
GAAGGAGCTA GTCTGCTGAA CCAGGAATAG TTTGAGATAT TGAAGTGTCA TTTTGNACA 180
TTTGAATACT TTGCAGGCTG GCTTTGTATA AACTTATCCT CNGGNNNCCT ATATGTTGTA 240
AATATNTAGN CCATAATTCC ATTATAANTA AANCCTATAA ACTATTCAA CNAANANCAA 300
ANAAACANANN 310

SEQ ID NO:6294

SEQUENCE LENGTH:258

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07456

SEQUENCE DESCRIPTION:

GATCCCCAAA AGTTTCCTTC TGCCCTTTTT CAATNCCTCC TTCCCGCCCN TCTGCTTTTG 60
TCATTATATA TTACTTTGCA TTTCTAGAA TTTCATATAA ATGGAATCAT ANAGTATCTA 120
CTCTTTGTTG TTGTCTGGCT TCTNTCACTC AGCATAATTA TTTTGAACT TATCTATGCT 180
GNTGTGGGTA TCAGTAGCTC ATTANCCNAT TGTGCTGAG TAGTNTTNA TTGTATGANN 240
GTGCTACACG TTATNCGN 258

SEQ ID NO:6295

SEQUENCE LENGTH:31

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07457

SEQUENCE DESCRIPTION:

GATCTTAATA AATAATACCT TACTGCTTAA A 31

SEQ ID NO:6296

SEQUENCE LENGTH:38

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear
 CLONE:HUMGS07458
 SEQUENCE DESCRIPTION:
 5 GATCTATGTA CAATTTTAAT AAAATCCTGT CCATGAAA 38

SEQ ID NO:6297
 SEQUENCE LENGTH:73
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07459
 SEQUENCE DESCRIPTION:
 15 GATCACCTAC TAAGANGTAC TTCCTACNTT TCATTCTTTC TTTATAAGAA TTAATTAANT 60
 TTTAATTGAC AAA 73

SEQ ID NO:6298
 SEQUENCE LENGTH:311
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07460
 SEQUENCE DESCRIPTION:
 25 GATCCACCCA CCTTGGCTTC CCAAAGTGCT GGGATNACAG GCATGAGCCA CCGCACCCGG 60
 CCCTATCTCA AGCTTCTTAA CTGCACCTGC TAAATCCCTT TTGCCATGTA AGGTAACGTA 120
 CCAACGGTTT CAGAGNTTAG GNTGTGGNCA TTTTGGGGGG CTGTTTTTCT GACTGCAATA 180
 CCTGGTNTCC AGTGAATCCA TGTTAGTGG AGGACAACAT TTGAGGCACT TACATGGATN 240
 CTAGGGGACA TNTATGTGAN TGTTAGGCAN AAAGTGCTGC CTNCANGNCA NGCAAATACC 300
 30 NATTNAATAT N 311

SEQ ID NO:6299
 SEQUENCE LENGTH:115
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07461
 SEQUENCE DESCRIPTION:
 40 GATCAAAAAA TAACTATAAG TATCCTCAGC CAAGTAGCCC ATGCCGCTGC TCTGCCTACG 60
 GNGTACTCAT CTTTATTCTT TTACTTCTT AATAAACTTG CTTTCAATGT ACAA 115

SEQ ID NO:6300
 SEQUENCE LENGTH:310
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07462
 SEQUENCE DESCRIPTION:
 50 GATCCTACTG CCTCGTCCTC TCAGAGTGCT GGGATTACAG NNGTGAGCCA CCACAGNAAG 60
 NCACAAATAG CGATTTTATC TACTACCACC AGAAATCTTC TGACATACTG TTTTACCGCA 120
 TGAATATTG TTTTCTTTAT AAATTCCTTA ACAAATCTCT GTTGTATAGA ATACTGTTGG 180

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TTAAATATA TCAGCAGCTA ATAGCTCATT TAAATAGAT TCTGAAAGTT CAAGNATTTC 240
AGGTTGCCAG NCACCNNTAT ACTCTACANA ATCAGANANG GCTCATTCTT GNTTCTNAC 300
ACAACAAANN 310

SEQ ID NO:6301
SEQUENCE LENGTH:209
SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07463.

SEQUENCE DESCRIPTION:

GATCAGGGAT GTCCAATCTT TTGGCTCCC TGAGCCACAT TGGAAGAAGA ATTGTCTTGG 60
GCCGCACATA AAATATGCTA AACTGACGA TAGCTGATGA GCTTAAAAA AAAAAANTTG 120
CANGAAAAAT CTCATGTTTT ANGAAGGTTT NCAGNTTGT GTTGGGCTGC ATTTAANGCT 180
GTNCTGGGCT GCNTTGGNCC CGNGGGCTN 209

SEQ ID NO:6302
SEQUENCE LENGTH:312
SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07464

SEQUENCE DESCRIPTION:

GATCATAGCT CTGGTAACT GTGGACTTGA TAGATTNAG GTACCTAGTT CAGAACTCCC 60
TAGTCACCAT CTCCAAGCCT GTCAACATCA CTGCATATTG GAGGAGATGA CAGTGGTAGG 120
ACCCAAGGAA GAGATGTGTG NCTGANTAGT CGTCACCATA TCTCCAAGCT TCCNGGCAAC 180
CAGTGGGAAA AGAAACATGC GAGNCTGTAG GAAGNGGGAA GCTCTTCCTT GGCANCTAGG 240
AGGNNTTAGC CATTCTCTTC CTTATGCAAN GATTGAGGAN TGCAACAATA TAAAGNACGT 300
GNAGTCCCA NN 312

SEQ ID NO:6303
SEQUENCE LENGTH:147
SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07465

SEQUENCE DESCRIPTION:

GATCCGCCCC CCTCAGCCTC CCAAAGTGTT GGGATTACAG GCGTGAGCAC CGCGCCTGAA 60
NCTGAAGGTT ACTTTTCTTT TCAGTGGTTT GGGGCAGGGT GTTACACAGC TTAATAGATT 120
AAGAAGAATA AAATGCTGAA TGCTAAA 147

SEQ ID NO:6304
SEQUENCE LENGTH:294
SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07466

SEQUENCE DESCRIPTION:

GATCATATTT CCTCAAAAAT AAGAGAAATA AGGTTTATAA GGAAACTTGC TGGGATTGTG 60

5 GTTGT TTTGT TTTCTCAGCA GCACAAACAA AACCAGAATT TAGCCTTTAG GACTGCTGAG 120
 TAAGCCAAAT TTAATGACT ACTGCTTTGT NCATGGGTAA GCCATGTNCT TTTCAAAATA 180
 AGTGCCACTA AAANCCACAT AATGCTTTGG TTTCTATGTG GCTAATANNT ATTTNGTCCT 240
 ATAGTTTATC TTATTTGTNA CTGATTTNCN TCTCTTGNAT GCCTCATATT ANT 294

10 SEQ ID NO:6305
 SEQUENCE LENGTH:294
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07467
 SEQUENCE DESCRIPTION:
 15 GATCCATTTCG CCCCAAGGGG ACAACGGGGT GTCAGGGGAG TNAGGGGTGG AGGTAGGGGC 60
 CAAAGAGGCT TACACGATAT CCCATACCTT TAATGCCTTT GGCCTTCCAT TCTNACTTCT 120
 CTNATGAGAT TATTGTCAAC CCTGCTTTCC CTGGTAGATA TTTGCCAGGC CCAATGCTTT 180
 AACCTTAAGC TGATATTTN CCTTTAGATG TCAATCTCGT TACCAGCAGC CTTTGACCC 240
 AACTNCAGNG NTCTATATTT NCGTAGNGGN TTTTCNCCA TGTGCATGGA TNT 294

20 SEQ ID NO:6306
 SEQUENCE LENGTH:129
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS07468
 SEQUENCE DESCRIPTION:
 GATCCCAGAG GCTGAGGGTT GGGGAGAGGA AGCCATCATC TCATTACCTC TGTCAGTGCA 60
 GGGGTGGCTN CTNCTCCCT TGTGCACTTG GGTGCTGTG ATTGTAAATA AACGTGATT 120
 30 CGTACCAAA 129

35 SEQ ID NO:6307
 SEQUENCE LENGTH:193
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07469
 SEQUENCE DESCRIPTION:
 40 GATCTCTTTG TNATTTAAGA AGTGGCTGGA TTGGAACCTT TAATATGCTA ATNTGGAAAA 60
 TTAATTACCT TTATGAAGGT GGTATTATAC AAATAAGCAC ACTAACCCT CGGAAGTTGT 120
 TTTACCTACT TTAAGGTTT TANTGGATTG CNCCTCTGTA ANCTATTCCT AANNTGTGTA 180
 TGANATATTT NAN 193

45 SEQ ID NO:6308
 SEQUENCE LENGTH:291
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07470
 50 SEQUENCE DESCRIPTION:
 GATCTTTCTT CCTCTCCAGC CCACAAGTGG CTCTAGTGAG ACGAGCATTT AAGCTGCAGG 60

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EP 0 679 716 A1

5 TGGTGGCACT CGGGTCTCAG AGCCACCAGA CGTGCAGGTG GGACAGTGGT TCCAGCTCCG 120
GCTGCCTGTN AGCGTCCCTC CCAGGTGCTA GCCCTGGCAG GCACAGCCCT NCCCTCATGC 180
CAGCAAAGGA AATGCTCTTG ATGAGAGGCT CGCTTTAAAG AAGCCCAAAG CGTGTGCTTA 240
TCCAAGGGGT TCAGCTATTC CTGACTTCAC GGCTTGCATG TTTGATTTTG N 291

SEQ ID NO:6309
SEQUENCE LENGTH:97
10 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07471
SEQUENCE DESCRIPTION:
15 GATCATATGT ATTTNTATGT TCTGTAAAGN ATTTGACTTA CTAGTTCTCA ATAAAAATTT 60
ATTAGGACTA TAAAAAATAA AAAAAANTTT TAATAAA 97

SEQ ID NO:6310
SEQUENCE LENGTH:147
20 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07472
SEQUENCE DESCRIPTION:
25 GATCTTATGT ACATACCCAT TTTAGCTTTC CCATGCATAC TTAAGTGCAC TTGCTTTATC 60
TCCTTGGGCA TTCTTACTTA GGATTCAATA GAAACATGTA CAGGGTAAAC AATTTTTTAA 120
AAATAAACT TCATGGAGTA TCTGAAA 147

SEQ ID NO:6311
30 SEQUENCE LENGTH:190
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07473
SEQUENCE DESCRIPTION:
35 GATCACATTT TGAATGCCT AAAAGACTTT ATTGTTCTAA TTATCCAGAT GTACCTTTGT 60
AAAATAGCTC TTTTATGAAT TAGCTGATAA GGCTGTATGT TTCTGGANCA AATATTGGGN 120
CACCTAAAC CTTCCGGTTT CCGGGGGCCN GGGAAATNGG AATNAGGTT TCNAATNTTA 180
AATNAGGNTN 190

SEQ ID NO:6312
SEQUENCE LENGTH:65
SEQUENCE TYPE:nucleic acid
45 TOPOLOGY:linear
CLONE:HUMGS07474
SEQUENCE DESCRIPTION:
GATCTGGATT TTTGCCAATT CCTATGAAAA ATAATCCCCA GAAATAAACT ACTAAGGAAA 60
ATAAA 65

50 SEQ ID NO:6313

55

SEQUENCE LENGTH:79
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07475
 SEQUENCE DESCRIPTION:
 GATCCATTGG TGTTTGGTTT GGGGTGTTTT TTAAGTTTTT TCTTTTATAT CATCCAGAAA 60
 TAAAGACACG TACACTAAA 79

SEQ ID NO:6314
 SEQUENCE LENGTH:204
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07476
 SEQUENCE DESCRIPTION:
 GATCCAATAA CCCAGTTAGG TATTATAGAT TAGTATTAA GTTTGCTGTA GATTGTGTGT 60
 GTGTGTGCTA AGTAAAAGTN CCATGATTCA CAATTTTAGT ATTAACCCAT GTTGCAAAAG 120
 CTGTTACTGG CCAACAGATT GTNATGATGT GTACCATATA ACACAAGCNT TANCTNGGCA 180
 TTANCGTTTG TNAAGCCANN TNTN 204

SEQ ID NO:6315
 SEQUENCE LENGTH:204
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07477
 SEQUENCE DESCRIPTION:
 GATCTCATCT CTGAAAATAA AATACCCCGT TAAAATTCTG GCTTTTGTG GTTAAGGTTA 60
 TTGAGAAGGT AATAGAAAAA GATATGATAC TACTTINCTC TCTTGGATTT GTAAAANTAG 120
 TGGAATTTTC ATAAGGATTT TGTATTTTCT ATTGCTGTAT AACATTATTA ATACAAACTT 180
 ATGGGCTTAA NANNCAAANN NNNN 204

SEQ ID NO:6316
 SEQUENCE LENGTH:218
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07478
 SEQUENCE DESCRIPTION:
 GATCACGAAG ATACATGTGT TTGACTGTTT AATTGAAAG TTTACATTTT TNATGCTTTG 60
 TGTGGTGTG TAATTTTGT ACTCTGGTG GCTAGTTTTT GTCAAATCTT TTTTGAATA 120
 TTGCTTAAAT GTNTTGATTT TATGATAGTG AAGCTTGAT TCAGTGTTTT GCCAATNAAT 180
 ATTATATGCT TGTAAATAANN GCAAAGANA AGCTTAAA 218

SEQ ID NO:6317
 SEQUENCE LENGTH:335
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS07479

SEQUENCE DESCRIPTION:

5 GATCCCAGTG GGAACCTTCA TGCCTTATTN ATTTCTAATG GGTAAAGGGG TTTTNTTACC 60
 AAGCATCCCT GACCTCCTGG AGACACCACC TGCTTTCCGG GCGGCACTGT GATGGGAGCT 120
 GGTGGCGACT GAGTCCTTCT GTACGTGCAA CTGGGAAACT TTTGTCTTT GAGGCTAGGC 180
 AGCTCCCTGC CCTCCGTGTG TGTCTGTTAT CTGGGGGAGA GGAGTGTGGG AAGGNNTTGG 240
 GGGAAGAGCT CCAGCCTGTN TTGNTCCCAA AGTCTCTNNAG TGGGAAGACC NGGGTNACCT 300
 10 TTNAGGGTN TANGGTGNGG GGAANTTNTT TTTN 335

SEQ ID NO:6318

SEQUENCE LENGTH:266

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS07480

SEQUENCE DESCRIPTION:

20 GATCTGCGTG TCAAACCTGTT CCATTTGTTT ATGTAAAGTG ATATTAATAA AGATATAAAC 60
 TATAACTGTC CGTTACTTTT GGCAAAAGAT ACAACCACAT AATGTATATA ATTCCTAGTT 120
 TCCATATTTA TCCGCATGTA AAGGGCCGGT TTATCCATGT TACAGCTCTT CAATATTTAT 180
 GGCTAGAAGA NCTCGTATGT ACACTTTAGT TTCCAGAACT GTTTGGTAAC CTTTCGTACC 240
 TNATTAGAGG ATTCTNAAAT CTCAAA 266

25 SEQ ID NO:6319

SEQUENCE LENGTH:313

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS07481

SEQUENCE DESCRIPTION:

35 GATCCACCCA NCTCGGCTTC CCAAAGTNCT GGGATTATAG GCGTGAGGCA CCGTGCCTTG 60
 CCGATAAATG GATTNAAGC TGAGTCTTTG AGGGAAGAGT AGGACCTGCA CTGAAACAAA 120
 AAGACAATTT GTGGGGTTGG GNGAGGAAC CAGATGTAAA CAATGGAAAG NTTTTGGTGT 180
 CACTTTGAGN NGGATTTTTT GGAAGGGTTT TGCAATGTCC ATTCTGTATC TTGGGTTANC 240
 ACTAGTAAGT CAAATGGNGA ATAATTTGAG TGCTACTCNT TCCCTNCAGT TNGGAAAAAN 300
 TTTTANCNG TTN 313

40 SEQ ID NO:6320

SEQUENCE LENGTH:144

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS07483

SEQUENCE DESCRIPTION:

50 GATCTGTGCT TTNCCCACT TTGCCCACT GTCTAAGGTG GCTTTNAACT GGAACCCAAG 60
 TGCAATAAAA GGTTGGTATT CGCTCCTGAA AAAAAAAAAA AAAAAAAAAA AAAAAANAAA 120
 AAAAAAANA AAATNAACNA AANN 144

55 SEQ ID NO:6321

SEQUENCE LENGTH:332

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07484

SEQUENCE DESCRIPTION:

GATCCCTCCT CACTTGCCCT GCCACTTTGC ATGGTTTAT TCTTAACCTG GTCCCCACG 60
 TGTNAAGTGT AGTGGCATCC ATTTCTAATG TATGCATTCA TCCAACAGAG TTATTTATTG 120
 GCTGGAGATG GAAAATCACA CCACCTGACA GGCCTTCTGG GCCTCCAAAG CCCATCCTTG 180
 GGGTTCCCCC TCCCTGTGTG AAATGTATTA TCACCAGCAG NCACTGCCGG GCCTCCCTCC 240
 CGGGGGCACT GCCTGAGGGC GAGTGTGGGN NTAGCATTAG CTGCTTNCN CNNNCCTGGN 300
 ACCNACTGTG GCCTGGAATN GNATAGTGGT NN 332

SEQ ID NO:6322

SEQUENCE LENGTH:376

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07485

SEQUENCE DESCRIPTION:

GATCTAATTA ATGTGAAGGA CTNGCCAGTC TGGGAAACCG CCTGCCACGT GGAAGAGCCA 60
 AACCCGACTC INTGCTGCCA CATGCCGTTT CCATGCCCGG CTGCTGGGCA CCTGGNAGAG 120
 CTTCCAGAAT CCTCGCAGAC AGCCAGAGC CTGCCGCTAC CCTCGGCCTG CCCACCACCA 180
 AGCAAGCAGC AAGCAAGATG GGGTTCTCAT CAGTTCTTCC TTCCACAAT GTAGGACCTT 240
 TCCTTTTACC TTTNCAATGG GATAAAATAG TTCAGAGTTC ATTAGTCATA TTTCAATTGGN 300
 CACAAGGAAT TCAANGGTTT TTTNANCAAT TTANCANTTC ACCCGTCTTA TGTNGTTTAA 360
 NANTNANANC NANTCN 376

SEQ ID NO:6323

SEQUENCE LENGTH:109

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07486

SEQUENCE DESCRIPTION:

GATCACAGAC ACCCAGTNCG TGGNCCCTGT CTTACCCCCA GCNAGCATCA TGTNAGTGGG 60
 CCTCGTTCCT CCCGGAGAAT CCCAGCGGGG CCTCANAGAT GCATCTGGN 109

SEQ ID NO:6324

SEQUENCE LENGTH:156

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07487

SEQUENCE DESCRIPTION:

GATCTTGGCT GCTTTTAACT AGAAGGTTGC TTTNATNAGC ATATTTATAC TGCTGAAGGA 60
 TGAGTGTNAA TTTAAATTAA CTTTGGCGTT TTGTAGAGAA AACTATTAC AAGATAAATN 120
 CCAAGTCTTT NCACCTGTCA GGCATGCATA TNTNAN 156

SEQ ID NO:6325

SEQUENCE LENGTH:229

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07488

SEQUENCE DESCRIPTION:

GATCCCAGTG AAGGGTTTTG TGTGTTTAGG CCTCATTTCT TTGTCTTTTT CCTACTCCGT 60
 TCCTGGCATT TCCTGATTTC NAGTGTATAC TCTGTAGTCT CAGTTCGTGT TTGATTCCAT 120
 TCCATGGAAA TAAAAAGTAT GTTGACATA CTGCCGAAGA ATTGTCTTGC AAGTAAAGGC 180
 TTCCCCCTTT ACTATAAGAC TATAANTAAA ACCTTATTN ATCCTTAAA 229

SEQ ID NO:6326

SEQUENCE LENGTH:190

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07489

SEQUENCE DESCRIPTION:

GATCTTACAC ATTCTGTGTA TAAAGACCTT AACTCCACAG GACGGACATT TTAGAGTTTA 60
 AATTATTAAG GCTATCATTG TTTTAGTAAT GTCATATTG CAAACTTTTT NAGTTTTGGC 120
 CTTTAATTTA AAANGCCTAA TTTTAANGTG CTGCCTGTNA GTAACCTTG ANTAANANCA 180
 ANTTATNACN 190

SEQ ID NO:6327

SEQUENCE LENGTH:355

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07490

SEQUENCE DESCRIPTION:

GATCTCAGCT CTTTNGGTGA CTGCATATGC AGTGGTGACC CAGCACAGTT TCCATCAATC 60
 TCGTCAAAAG ACTTGAGGTT GTTGGGCACA GTATTCAGA TGACCACAGT TAAAAGCTG 120
 GGTATACACA ATTAACANCC AANGTGATAT GCATTTATAC CTTTCCCTTT ATGNCCAATT 180
 TCTTTATAAA TATGGCTCAT CTGCTCATAA CTGTTAGNCC CACCCAAGTG TCATTAGTAA 240
 CCTGAGTGTT TATGCTTGA AACTGTATG TTGTTATTGT CTGATTGAT NATATTCNGT 300
 GNTGTTTAGN GAATGCNGTN GTGTTTNANT GNATCACAAA NAATTNCCCC ATTN 355

SEQ ID NO:6328

SEQUENCE LENGTH:298

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07491

SEQUENCE DESCRIPTION:

GATCATTTGA ATCTCAGTCT TCAGCCTGCA CTGATTTGTA GCCTGCACTG TCTTACTGAT 60
 TTACAAAGTG AAATCACTGA GAAATGTCTT TAGTTCAGTG AGAAGAAACC AGAACAATTG 120
 TTCCTAGTGT TGTGTTGTTT TTTTAAAGCA AATTACTTAC TGTATTTTNA TGGCAGGNGG 180
 GNGAAAAAGT GTTACAACGG TTTCTAATGA AGTCCGGTAT TTANNTGGA AATGACTAAT 240

GTGTTTAGTA GTGCCAAANT AAACCATTAN NTGNTTGTCC TTTGCCAAAN ANANNANN 298

5 SEQ ID NO:6329
SEQUENCE LENGTH:225
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07492

10 SEQUENCE DESCRIPTION:
GATCACAAG AAAAAGCATT TTAAAAAGT TGGCAAACGC TGAAACGCAC TGTGGTATGA 60
GGCGCATTGC ATTTCCATAG CACTGAAGTA CCAGTTTCCA TTCCTGGGCT GNGATTGTTT 120
TTCCCGTGGT TGTATTGTTC TGATTAAAG ACCACCAGNG TAACTGNTTT TTNTTTGTTT 180
15 GTTNTCTGGT GGGGTNNCCA CCAATAAAA CTNNTANAAN CCCGN 225

SEQ ID NO:6330
SEQUENCE LENGTH:229
SEQUENCE TYPE:nucleic acid
20 TOPOLOGY:linear
CLONE:HUMGS07493

SEQUENCE DESCRIPTION:
GATCGACTTC GATAAAATAC TTTTGCTAA TCAAAATTAG AGTGTGTTT TGTCTGTGT 60
25 AAAATAGAAT TAATGTATCT NGCTAGTAAG GGCACGTAGA GCATTAGAG TTGTCTTTCA 120
GCATTCAATC AGGCTGAGCT GAATGTAGTG ATGTTTACAT TGTTTACATT CTTTGTACTG 180
TCTTCTGCT CAGACTCTAC TGCTTTTAAAT AAAANTTNAT TTTTGTAAT 229

SEQ ID NO:6331
30 SEQUENCE LENGTH:345
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07494

35 SEQUENCE DESCRIPTION:
GATCTAAAGA CCCGATAGGT GCAGAACCCA TCTGGACACG GAGACCAGGA ATGGAGTTCC 60
ATGGAGGCCT GGCTGGCACT GCACCCGGGC ATGAGGACAC ATCCAGTAAG AAGACCTGCC 120
TCAAGAGGTG CACTGCGGTG ACCAGTGGAG GTNACTGGTT GGAGCCTGGA ATTGGAAGCA 180
GATTCCAAGC TCTGGTGGAC AAATCTCCA GGCCTGGTGG GATTACAGC TGGGGCAGNC 240
40 CTCATCCTGG CTGCNTGGCC ACAGNCCCN GCTNTTNCCT ACTTGGTGGT AGGGCCGATG 300
CCTTNTTGG ANGAGNTGGG NTTNTTTGG TCNCGACTGN GTANN 345

SEQ ID NO:6332
45 SEQUENCE LENGTH:219
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07495

SEQUENCE DESCRIPTION:
50 GATCACAAG AAAANGCATT TTAAAAAGT TGGCAAACGC TGAAACGCAC TGTGGTATGA 60
AGCGCATTGC ATTTCCATAG CACTCNAGTA CCAGTTTCCA TTCCTGGGCT GAGNTTGTTC 120

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EP 0 679 716 A1

TTCCCGTGGT TGNATTGTNC TGATTTCACG TACNCCAGAG TAACTGNTTT TTTTTTGTTT 180
GTTTTCTTGT GGAGTTAACA CCANNTAAAA NTTGTAAAN 219

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SEQ ID NO:6333
SEQUENCE LENGTH:60
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS07496
SEQUENCE DESCRIPTION:
GATCTATTTC TCAGCTTTCA CTTATGTGAG CCAATAAATT CCTTTTTTGT TGAAGGC AAA 60

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SEQ ID NO:6334
SEQUENCE LENGTH:154
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS07497
SEQUENCE DESCRIPTION:
GATCTACCCC CCCATCAACG TGCCCCCTCG ACTCCCGGTG CTGCGGAAGA ACTGAAGGTT 60
GCNATGCCTT ACTCTGACGG GAGCATCTGT ATTTTAATGT TAAAAGCCCA CAAAATAAAA 120
ATAAAAAGTA ACTGAGATGA ATTTAATAAA CAAA 154

25

SEQ ID NO:6335
SEQUENCE LENGTH:91
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

30

CLONE:HUMGS07498
SEQUENCE DESCRIPTION:
GATCTATGCT GTATGCACAG ATTGCATAAT GGCTTTTTTN CTTAACATTA TGTGATAAGC 60
ATTNCTGA ATAANCTTAT TTTAAAATAA A 91

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SEQ ID NO:6336
SEQUENCE LENGTH:181
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS07499
SEQUENCE DESCRIPTION:
GATCAGGTGT GGCCAATNAA ATTNAAGAGG AAGTCTACCT GGGNTTCTGG GAAAGCTTTT 60
CCAATAAAAG ACACAGGCAT GGCTAACACC TCCCTGGGCC TCTTCTTCCT ACCTTGATTG 120
AGGGTGTGAT GCCTGGAGCC ACAGCAGCCA CTTTGNNNNC ATGACAAAAA GGCCAAGNGN 180
N 181

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SEQ ID NO:6337
SEQUENCE LENGTH:77
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS07500

SEQUENCE DESCRIPTION:

5 GATCACCAAA ACCTGATGTT TTAAATNTGC TTTCTTTTGG AAATTTATGT TTTCAAATAA 60
AATCTCCCTA AAGCAAA 77

SEQ ID NO:6338

SEQUENCE LENGTH:265

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07501

SEQUENCE DESCRIPTION:

15 GATCCTATGA GTGTAGTTAA TGACTGTTNG TAAGTCAGTA GAGTAAAATN CTGTGTCCAC 60
GGGGTGTAC AGCCTACCA TACCCTGTTG AGGTGTGAAA TGCCCCGTCA GAAATTAAAT 120
ACAAACTTAA ATGTGCCTAT TGGTGTCTAA NCTTCATACA ATGTAAGGNC AGATTCCTTT 180
TNGGNNTACT GGGTGTCTGTC GCCAGGTTTG NTAGTTAGAC TAAAAANCTT GANATTCACT 240
TTTGGGGGG NGGNTATAC TGANN 265

20

SEQ ID NO:6339

SEQUENCE LENGTH:161

SEQUENCE TYPE:nucleic acid

25 TOPOLOGY:linear

CLONE:HUMGS07502

SEQUENCE DESCRIPTION:

GATCTATTGG AAAACAACAT GGAATGGAAT TCTGGAAATN ATTATTCATT GAAGAATGCA 60
GTGGCCAAGA AAATATCAAA TGTAGATTGT NAACGCTTGA GAATCATGGC TATGGTTTCT 120
30 NATGTCTTGG TAACANGCTG TTATCTTTNN NGACATTTN N 161

30

SEQ ID NO:6340

SEQUENCE LENGTH:272

35 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07503

SEQUENCE DESCRIPTION:

40 GATCTAGTTA TAAAATACAT GAAAAGGCTG ATGCAGCAAT CGGTGGAATC GGTTTGGAAT 60
ATGGCATTGG ANTTTATTCT TGACAATGTC CAGGTGGTTT TANAACAAAC TTATGGAAGC 120
ACATTAAAAG TTACATAANT ATTACCAGAG AGCCTGATGC TCTCTGATAG CTGTGCCATA 180
AGTNCTTGTG AGGTATTTNC AAAGTGCATG ATAGTAATGC TCGGNGTTTT TATAATTNNN 240
NNTTCCTTTT AAAGCAAGTN TTTTGTACAT NN 272

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SEQ ID NO:6341

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

50 CLONE:HUMGS07504

SEQUENCE DESCRIPTION:

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5 GATCAGTGTG GAAAAGCCTT CANTCAGAAA GGAAGCTTAA TTNTNCACAT CAGAGTCCAC 60
 ACAGGCCTGA AGCCCTATGC CTGTNCCCAG TGCAGGAAGA GTTCCACAC CAGGGGGAAT 120
 TGTATTCTGC ATGGCAAAAT CCACACAGGA GAGACACCCT ATCTNTNCGG CCAGTNTGGA 180
 AAAAGCTTCA CCCAGAGAGG GAGTCTGGCT GTGCACCAGC GAAGTCTTCA CAGAGGCTCA 240
 CCCTTTGACC ACTTTCCTGA AGAGAAGTTC TCTTTATGAA TTAAGGAGTA CAAAATCCTC 300
 TGAGNTTGAN GCAACCCTAT CCAAGTTTCT ANTGGGANTG GTTGN 345

10 SEQ ID NO:6342
 SEQUENCE LENGTH:212
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07505

15 SEQUENCE DESCRIPTION:
 GATCTGTCTG CCCATCTGGC CCAGGGGGTC CGAGAAGGGA AGCCTTGGGC AAGAGGNGAC 60
 CAGTTGCAAT ACTGTACTTC CTGGTCAGTG GCCAGAGGAT GCGTGCAATA GCAGAGGCCA 120
 GGTNACCCCT TCAGCCTTGG CCTCTGCCNC TCCATTGGCC CTCCCTNCCT GNTNCTNCNT 180
 20 GGTGTNGGTC AGTCCTTTTC TAAAGNTGTC CN 212

25 SEQ ID NO:6343
 SEQUENCE LENGTH:316
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07506

30 SEQUENCE DESCRIPTION:
 GATCTTCGTA TAATTGGCCA CTAATGTGAG AGTTCACTAC TAGGCAGAAA CTATTATGGA 60
 CAGTNAATA ATGACTTTAA TCTACCACG TGAGTTTAT GCAGTCTTTC CTGTCTAGCC 120
 CTGCCTCTN CCTGCCCATG TNATTGCGGT GCAGTAGTTT CTGTNGTATA ATAGTGTGGA 180
 CAGCAGCTCA GAAAAGGAGG GAATGCTACT GATAATTTGT AGATAATNN NTTTANGACT 240
 TAGGGGAACC ATTGACCTTT GANATTTTNA TTAGAAAATN ATTTGTNCAG AATCAGACTC 300
 35 CATTATTTNC CATATN 316

40 SEQ ID NO:6344
 SEQUENCE LENGTH:184
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07507

45 SEQUENCE DESCRIPTION:
 GATCAAAAGG AATGATGTAT TTTTAACGTG TTGGCCAAAG TCACTGGATA AAATGAGAAT 60
 TGTATATTTN TAATTCATTT TGCAATNCA GAAAGTTGGT CCAGATATAT GTCACAGANC 120
 TTTNCACITG TATACTACTC TTACAATGGA AAANNNTCCC GAAANCTGTA TACTTCTGAT 180
 TANN 184

50 SEQ ID NO:6345
 SEQUENCE LENGTH:361
 SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS07508

SEQUENCE DESCRIPTION:

5 GATCAGACCA TGCAGGTAAG GTGAACCAGC TGCACGGACC AGGTTCCCGC AAAACATTGC 60
 CAGCTAGTGA GGCATAATTT GCTCAAAGTA TAGAAACAGC CCACCTGTGC CCACTTTGAC 120
 CATTGGTGAG GATAGATATA AAATCACTTC TTCCAACGAA GCCTAGGTGA AAATCTATTT 180
 ATAAATGGAC CACAACCTCG GGGGTGCGTT TTTGTGCTGT GACTTCCTAA TTATTGCTAA 240
 10 AGAACTACTG TTTAGTTGGT AATGGTGTA AATTACATTC AGCTCCTTCT TGTCATATAA 300
 ANAGGAATTT GGNNGGTGTC GGTTTAAANA TTTTATTNCA CCTGTACCAT TTNGTCACTC 360
 N 361

SEQ ID NO:6346

SEQUENCE LENGTH:266

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07509

SEQUENCE DESCRIPTION:

20 GATCTGTTGT TATACCAATT GAATTAATTG AGTTNNTTGT TGGACATTG GGTGGGTTNC 60
 AAGTCTTTGC TATTGTGAAT AGTCAATGGC AGTATTTTNA ATGTTGGTCT ATTATTGTAC 120
 TAAATAGTTG TTTATATTGA GCTTGGGAAA GCTTTCGTGT CCAATAGAAA TAGCATATTA 180
 25 AAATAAATGC CACTGTATAT NAATGANAGT GTTGATTATA TACCCTAGAN ACTTAANCCT 240
 TAAGGNAGTT ATGTNTNAAT TNNCTN 266

SEQ ID NO:6347

SEQUENCE LENGTH:355

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07510

SEQUENCE DESCRIPTION:

35 GATCAACTCC AGTTTCCAAT GTCTATGTGT CTATGTGTGT ATGTNCCATA CATATGTATT 60
 CACATGAAGA CCGCATGGC CAAGTNCTGC TGGAGGAGCA CTCAAGTGTG ACGAGCAGGG 120
 CCACTGGACC CTGCAGGGCT GTGGTGTATA TAGTGCAGCT TTGGAGGTGG AACTCTATTT 180
 NAACACTTTT CTATGGAGCC TTCCGAGTCC CAGGTTTNA CTTGAGGCTG TCTGCTGGA 240
 TGGCGGTTTT NAGACCTCCA TTAACATCCC TACCCAGCAT TCTGTACTTC GGGGGGNCTT 300
 40 CTATCTTGTN ATAAACTTN TTACCAAGTG GAACATCGGT TACCCACCTT TTGTN 355

SEQ ID NO:6348

SEQUENCE LENGTH:358

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07511

SEQUENCE DESCRIPTION:

50 GATCGTCCTG CCTTTGTAAG GCAGATTGG AGGGGAGGCC CAACAGCAAA GATGGAGAGG 60
 AAAAAGGGAG TGGAGCTTGT GGAACAGGAA GCCCCCAGG GTCTGTCCAG GGCTGCTCAG 120
 CTCTTNAGGA TGTTGTGGAT GGTGAAGTGG TGATGGTGTC CTGGAGCAGC CCAGCACCTC 180

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EP 0 679 716 A1

5 CACTGGGGCC AGAAGGTCTT GACTTCGGAC ACTTACCCCT GTTTCACAAG TGTTTATAAA 240
GCTGTTTTTN CITTINGTTTT TCTCCTAAAT AATCCTTACT TGGTAGTTAT NAATCCTTCT 300
TTGGGGGAGG GACAGGGGCT TTAANCTG GAGGAAAATT AATTTNNAT TAANGGAN 358

10 SEQ ID NO:6349
SEQUENCE LENGTH:106
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07512
SEQUENCE DESCRIPTION:
15 GATCATGGAA GCTCCGTGCC CCTTCTCCCA TTCCTTGCCC TAAGTATCGC CTCCATCTGG 60
TTGTTGTCT GTATCCTTA TTAATAACA AGTAAATNTA AGTAAA 106

20 SEQ ID NO:6350
SEQUENCE LENGTH:135
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07513
SEQUENCE DESCRIPTION:
25 GATCAGTGCC AGAAGTCATT GTATTCAAAG AAGAATAAGC AAGAAAGAAA AGAAGGAAGG 60
AAGAGAGGTA GACAGATACA AGATGAAATC CTGTCAAAAA ATGGAAGGAA ANCCAGAAAN 120
TGAGAGTNNN CCANN 135

30 SEQ ID NO:6351
SEQUENCE LENGTH:156
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07514
SEQUENCE DESCRIPTION:
35 GATCAGGAAC ATGGCAAGGA ACACCACACC ACTGCTTTN AGCATTGTAC TGGAAGTNCT 60
ATATAATGCA GTGTTAAAA AAAAGAAAT AAAATGCACA TAGATTGAA AGGNTGAAAT 120
ACAACTTNT NCACAAGTAA CATGATTGTN TTTNTN 156

40 SEQ ID NO:6352
SEQUENCE LENGTH:134
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07515
SEQUENCE DESCRIPTION:
45 GATCAAAAAG AACTATGAG TAACAAGCTA TAACATAGTT CACCACAATG GGACCCCCC 60
CCCTTTTCCT NACCCTACAG TTAGTAATAT TNCAATTAAA NTANCTATAT TCTNCTATAT 120
TTNCCTGTT AAAN 134

50 SEQ ID NO:6353
SEQUENCE LENGTH:107

55

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07516

SEQUENCE DESCRIPTION:

GATCCACATA TCTCTAACAC CATTAATATG AATCATTGGA AATATTTTAC TTATTAACAA 60
AGTCTCTTGT GTTAAGCTAC AGATAAAGCT TTTGTGGTAG TGTCAAA 107

SEQ ID NO:6354

SEQUENCE LENGTH:350

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07517

SEQUENCE DESCRIPTION:

GATCCAGAGA CATGTGGCAG CAGGCATGGC TTCCCCTTGG CCTCTCTGTA CACTGCCCCA 60
GGACTGTCAT TTTGGCATCT GCAAAGGAAT CACTTTAGAA AGCCAGCACC TGGTTGATGT 120
GTATTCATAC TGACATTAGA TTGATGTGCA CTGCAGGGGN AANNGCGGTA GCTGNCACAG 180
NAAAAGGATG TTTTGATAGG AATAATTTTC TAGTATGTCT TGAAACATGT TCATCTGGAA 240
GTATTTTCCT CCAAAGTAAT GTAGCATGAT TTNCAAGGA TTGTTAACAT GCCTGGGATT 300
GGGAAAGNTA GGCCTAANGT TGTGCCAAAC TATATCAATA AATTCCTTN 350

SEQ ID NO:6355

SEQUENCE LENGTH:137

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07518

SEQUENCE DESCRIPTION:

GATCCACATT GCATGCACCC AGAATTTTAT ATTCCCAGG AATATAGCAG AGGAATATTC 60
TGGTATCATA TTTATCTTTC ACCATCTCAA AAATTACAGT TTTAAAACT CAAATAAAAT 120
TCATCACTAT AACCAA 137

SEQ ID NO:6356

SEQUENCE LENGTH:135

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07520

SEQUENCE DESCRIPTION:

GATCCAACCC TTGCTAGGTA TCATAGTTTA GGCCCATTTA CCTTCCCCTG TACTGGCAGT 60
TCAGCCGCTT ACATGCACTC ACCCTGTTTG TGGCTATTTT AAATTCATAT TATTAACAAA 120
CAAAAAAAC CCAA 135

SEQ ID NO:6357

SEQUENCE LENGTH:69

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07521

SEQUENCE DESCRIPTION:

GATCCTGTCT CAAAAAAAC ATAACCGACA TCCAGAAGTG AAATAAATTA GCCAGTGTCA 60
 CAAAGTAAA 69

SEQ ID NO:6358

SEQUENCE LENGTH:200

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07522

SEQUENCE DESCRIPTION:

GATCTTAAAA NTACAAAATG TAGACAATTA ATATTCAGGT GCTTACTTTT GAGCCAATAT 60
 GTAGCAGAGA AAAACTAATT NNTCTGTGGC AGTATTTTNA TTTTNGGCAT TTCCAGGAA 120
 TTAAGTGAAG ACTAAACTC ATATGTGANG TGTAATGATA CTANGGNAAG AGTCAGTAAA 180
 ATTNCCATTT GGCTACCAAA 200

SEQ ID NO:6359

SEQUENCE LENGTH:115

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07523

SEQUENCE DESCRIPTION:

GATCTTGTAAT ATATATAAAT NTAAATAANA TTAAANCTGT AATTCCTTTN CCCTCCAAAG 60
 GCTTTTNTGT ACATGGCGCT GCATTTGGCT ATTCCTTTG GAAATAAATA ATGTN 115

SEQ ID NO:6360

SEQUENCE LENGTH:319

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07524

SEQUENCE DESCRIPTION:

GATCTTCCCA CCATCACAAA TGAATTTAAA GATGAAAAGA AACTCAGTTG CTCATACAAC 60
 TGCATTTTTC CTGTCTATTA TGGGAAACAT CAGACGTTCT GAGTAAGATA TATCTCATGG 120
 CATTAGTTAA TATAACTNAT ATTGTTTAA TCAATGGTATT ACATGCAATT NATATCAGAT 180
 AAAAGCAGAA CACATTTTGT TNCTGCCTCT CTTAAATGCT GAATGTAGCT GTNATGTATA 240
 ANTCCATTTA GTTTTATGTN CTAAAGGNCT ATTTGTGCAC TCCAGNITTN CAGTAAANTA 300
 GNATTCCTAG TAAANCNNN 319

SEQ ID NO:6361

SEQUENCE LENGTH:141

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07525

SEQUENCE DESCRIPTION:

GATCCTAATT TNNTACAGAA TCCCGATGGA ACCCTGTGCT GTTNANACAT CAGTGTGTAA 60
 AACTCTCTGT GTCCCTGTTG GGCTGTCCAG ACAGTCTGGA CTCTGTGAAA TTAAANATTT 120

AATTAAAGGA AACAAACCAA A

141

SEQ ID NO:6362

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07526

SEQUENCE DESCRIPTION:

GATCTTTTGC TTAGACGTTA ACTTGATGCA TCATTGGAAA GNTGTTTCTC TCATCTCTGT 60
 CCTAAGGCTT GATAAAGTCA TTAATTTGT NTTCTTTTGA CTAAT 105

SEQ ID NO:6363

SEQUENCE LENGTH:392

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07527

SEQUENCE DESCRIPTION:

GATCTCAGGC CAGCCGTTTG TNCTCCTAGC AGGGTTGCTG TGCTGGCCAC ACGGAGAGGC 60
 CCTAGAGAGC CTCATGGATT GTAACATAAG AAGAAACGGT TCCTTTTGT TTTTAAAAA 120
 ATNATTTTAA AATACCGTTT TTAACACCGT TCTCTCGGTA CTTTTTTTAA GCTAAGTCAG 180
 CATTGTCTTC CAGTGTTAAA GGCATCCCTC ACCTCTGCAT TGACCTTACG NATCCATGCC 240
 AAGGAATGGG ATTTCCATCC CTGNGGCCAG TCAGTTAGGG TGCAATTGT NCCTATTGA 300
 NTTTTTAATG NAATCTGATG NGGTGAGNTC AGGTTTANAA NTCTTCAAAN GNNCGGTNA 360
 TTGTACCAGG NNTGNTATGG ATTAATCCNT GN 392

SEQ ID NO:6364

SEQUENCE LENGTH:205

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07528

SEQUENCE DESCRIPTION:

GATCTCCATT CCTTGGGCTC CGCGTCCGAN TTCATGGTGC GCCGCTGTGC TGGGAGCTGC 60
 AGTGGGAATN TNTGGGACAC CTTGACCAAA GGGGAGCTTT GTNTCGTGTG TTTTGAAAAA 120
 GGCTTAATGA AGAGAATGTT GTTCATTCTT AGTAGTATAG TTTGCAATTC TTAATGGCAA 180
 ATAATAAGTT TCAGTAGAAA ACAA 205

SEQ ID NO:6365

SEQUENCE LENGTH:343

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07529

SEQUENCE DESCRIPTION:

GATCCTTTTC TACCCACTTT CCTATGGAGG ATTCCAAGTN ACCACTTCTC TCACCGGCTT 60
 CTACCAAGGT CCAGGACTAA GGCCTTTTNN TCCATAGCCT CAACATTTTG GNAATCTTCC 120
 CTTAATCACC CTTGCTCCTC CTGGGTGCCT GGAAGATGGA CTGGCAGAGA CCTCTTTGTT 180

EP 0 679 716 A1

GCGTTTTGTG CTTTGATGCC AGGAATGCCG CCTAGTTTAT GTCCCCGGTG GGGCACACAG 240
CGGGGGGGCG CAGGTTTTCC TTGTNCCNCA GCTGCTCTGC CCNTTNCNN TTATTNCCTG 300
ACTTCAGGNC TTGAAACNNT TCCNGTGGTG TAAATAAAAA ANN 343

SEQ ID NO:6366
SEQUENCE LENGTH:332
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07530

SEQUENCE DESCRIPTION:
GATCTGTACT TTGAAACAC ATTGAGTCCC TTAAACCAGG AAGAACTCTT CTGGGCGACC 60
GTCCACATGT TCCTTTCCT TCCTAATGAC GTGAAGTCTG AACTCTACAG AGCCTCTGTA 120
CCTTCTCTTT AAAGGCAGCT CTGCTCACCA GGAATTGGCT ATAATNCTTT CCAACTCTCT 180
CATCATTTTA ATAGCTATCA GCAAATGTAG AGGTGTAGGG GATTTTGTA CATCTGATGT 240
ATGTAAAAAG ANATATTCAA GTTGAAGGC AGNCTAGAAC TGNCTGNGA GATGTTTGAT 300
GTACACATTA AANNTAAGGA ATTTACATA AA 332

SEQ ID NO:6367
SEQUENCE LENGTH:258
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07531

SEQUENCE DESCRIPTION:
GATCAAGAAT GGCAAGAAAA TCACAGCCTT TGTACCCAAT GACGGTTGCT TGANCTTTAT 60
TGAGGAAAAT NATGAAGTNC TGGTTGCTGG ATTTGTCGC AAAGGTCATG CTGTTGGTGA 120
TATTCCTGGA GTCCGCTTTA AGGTTGTCAA AGTAGCCAAT GTTCTCTTT TGGGCCTATA 180
CAAAGGCAAG ANGGNAAGGC CNAGNTCACC ATCCCCTAAA TTTTAAAAG NAGTTTCACA 240
AAGTNTGCA TATTTNNN 258

SEQ ID NO:6368
SEQUENCE LENGTH:148
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07532

SEQUENCE DESCRIPTION:
GATCTGTACA TTATTGCATG TGGCAGTAGT GTATTTNCAT TTTGCATACT ATTCCATTTT 60
AAGAATATAT TGCTGTTTAT CCTATTGATG GATATTTGTA TTTNTNATAC TTTTGTGTAA 120
TAATAAAAAT ACTGCTGTAA ATATTAAA 148

SEQ ID NO:6369
SEQUENCE LENGTH:344
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07533

SEQUENCE DESCRIPTION:

5 GATCTGATTC AAAGATAACC ATTACCCATC TTTCCAAGGC CCATCTAAGT AGTACAGGTG 60
 TTTTGTCTTT TAAAATTGAC AGTCCTCTTT GCATTAAGTC TGTAGGGGGG GAATACAATA 120
 TAAATATTCT NCTGTGTAAA CGTAAGAATG TGCTGTAATC TCCAGTTCCT TTGGATACCT 180
 CAGAATTCTA TTTATTGTAT ACACTTATAC TGTTTATNCA ATGTTTTGGG TCTTACTCTT 240
 TTTGACGTAT TCTCATCAGT CTTCACCTA GTTTGCAAGC NCTGAAGTTA TTGATTACGT 300
 CTAATGTGCA ATTNGGCTTC TATTAANTGC TATTTNATGG TTCN 344

10 SEQ ID NO:6370
 SEQUENCE LENGTH:292
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07534

15 SEQUENCE DESCRIPTION:
 GATCTNTGTC ACACCCAAGC TGGTGGAGGA GCACCTCAA AGTGCCANT ATANGAAACC 60
 ACCCATCACA GTGGACTCCG TCTGCCTCAA GTGGGCACCC CCCAAGCACA AGCANGTCAA 120
 GCTCTCCAAA GAAGTGAGCA GCCTGGCCCC TGCTGTCGGA CCTGAGCCTC CTGGCTNNGG 180
 20 GNCTGTAAAT ATGTATAGAC CTGTTTTGTC ATTTTTTTAA TAAAGTCAGT TCTGGTGGCC 240
 CTGNGACTTT GGNAGGGNAA GGGGGAGGCC ANAAANANTA AANGGGGGGA AN 292

25 SEQ ID NO:6371
 SEQUENCE LENGTH:139
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07535

30 SEQUENCE DESCRIPTION:
 GATCCACCTG CCTTGGCCTC CCAGAGTGCT GGGATTACAG TCGTGANACC GCACCTGGCC 60
 AATTTTAATA TTTTAACTGA AGATGTTTCA TATAATATTC AGCAATATAA AAATTACTAA 120
 TGACTTTTTT GGTAAAGAAA 139

35 SEQ ID NO:6372
 SEQUENCE LENGTH:105
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07536

40 SEQUENCE DESCRIPTION:
 GATCCTAACC GNAGGCCCAT CAATTGCAAC ATAGAGATAC ACATGAAAAG AGGGTTTAAC 60
 AGACCCTCTC TNGGACACAN CTGTTTCTCC TTTGAGCANT ANTTN 105

45 SEQ ID NO:6373
 SEQUENCE LENGTH:151
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07537

50 SEQUENCE DESCRIPTION:
 GATCCAGACC AGCGTCTGGC CCAAGCCATT CANAGGTGGG GTCTGGGGNC TGGGGGTTTT 60

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EP 0 679 716 A1

GTNTCGTGAC ACTTTGTCTC AAGGAGATTC CACATAGGGA TTTNAATTAG GCTTTCTNCT 120
TGCNAAGGCC TGTGGCATCT AANCCTGGCT N 151

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SEQ ID NO:6374
SEQUENCE LENGTH:288
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS07538

SEQUENCE DESCRIPTION:

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GATCTNAGGC AGATAAAGAA ACTTCAACTT ATTNAGAGA GGAGGTAGGG GATGGGGAGG 60
TCACAGAGAA CTCTGGGTTT CTNCAGTTA CTATGCCACA GCACCATATT TCCGGGTATG 120
AGTTCTGAGC CCCACAATGG CCATAAGCAC TTAACCACAG ACCTAGTGAC GTATGTATAA 180
NATATACT AGATTCCAAA GNCTTAGTAT ACAAAGACC ATAAATATC TTATTTGTAA 240
TTGTTAANA CTGATTACAT GTTAANNNTGN TAATNTTTN NNTATANN 288

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SEQ ID NO:6375
SEQUENCE LENGTH:319
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07539

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SEQUENCE DESCRIPTION:

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GATCCTCTAC CCCCGGAATG AGGACACACT TCAGGACCCA GCCCCACTGG AGTNTGGCCA 60
AGNATTCTCC CAGCAGGAAA ATGGCCATTN CATGGACACC AATGANTGCN TCCAGTTCCC 120
ATTGCTNTGC CCTCGAGACA AGCCCGTATG TNTCAACACC TATGGAAGCT ACAGGTGCCG 180
GACCAACAAG AAGTGCAGTC GGGGCTACGA GCCCAACGAG GNTGGCACAG CCTGCNTGGC 240
TCAAGTGGCC TTTTLAGGTG GGTATTCTTC AGCCGTCTCT AGAATCTTTN AGCCTCTTTN 300
TNGGGCCTCA NATCTTGN 319

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SEQ ID NO:6376
SEQUENCE LENGTH:166
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07540

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SEQUENCE DESCRIPTION:

GATCTCTGTA TGNTCTGTGT GTTCTGTGT CCTGGAATTG GATGCGTGGG ACTCGTCTG 60
TCCGCGGAGT GCACTCTTTT TTNCAGTGTG GCCACATAT CTTGTAAATN TTTGCTGAAG 120
AGTTGTGTCT ATATATAGAG AAAATATATA TAANCAGAGA AATAAA 166

45

SEQ ID NO:6377
SEQUENCE LENGTH:158
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07541

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SEQUENCE DESCRIPTION:

GATCCAATGC CCTAACTTT ACATAAGAGT TCCGGCAAGG CTGTGACTTC AACTTGTGTT 60

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EP 0 679 716 A1

GTAAGTCAA CCTGCAGAAT CAAATTTAGG GTTATGTTTT CAGTTGTATC AGTCTTACGG 120
CTACAAGATA TTAAGAGAT TCTTTCCTGG TTATCAAA 158

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SEQ ID NO:6378
SEQUENCE LENGTH:323
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07543

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SEQUENCE DESCRIPTION:
GATCTCACCT TGTGTGCAT CATAGAATTC AACTGGACT AAAACCTTTT AAGTGTAAGG 60
ATTGTGAAA ATGTTTTAGT CGAAGCTCTC ACCTTTATTC ACATCAAAGA ACCCACACTG 120
GAGAGAAACC ATATGAGTGT CATGATTGTG GAAAATCTTT CAGCCAGAGT TCTGCCCTTA 180
TTGTGCATCA GAGGATACAC ACTGGNGAGA AACCATATGA NTGCTGTGAG TGTGGGAAAG 240
CCTTCATCCG GNAGGATTGN CCTCATTANG CACCNGNGAT TTCATGTTNG NGNAGAGGCC 300
TATANTTGTA ATCAATGGTN ANN 323

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SEQ ID NO:6379
SEQUENCE LENGTH:318
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07544

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SEQUENCE DESCRIPTION:
GATCCTTCTA ATCTCCTGGA CCCTCCAGGG CACTCTGGTC CCTATTCCCC AGCTCCTAGG 60
CAGCTGAGCC GGGTCCCTTA GGGGAGGTGA CCAGGAGCTT TGGTGCAGGG AGCTCTTNGT 120
GGGGCAAAGG GCTGGACCCC TGCCAGGTCT GTGGACATGG TTATATGCCC GGNGAGGGG 180
GGTACAGGGC CCCAGGGATG GCCCCAATC CCACCTCTGT TTATTCTGTA AACTGCAACC 240
TATAATAAC CTTAGCATT CCTATTGTA ACAAATNG NNTTTNTGNA AATAAATTAT 300
ATTCCTAGT CTAACAAA 318

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SEQ ID NO:6380
SEQUENCE LENGTH:77
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07545

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SEQUENCE DESCRIPTION:
GATCTTCTCT GCCTTTATTG TTTTGCCATT TCACTAAAAT ATGTTTAGAG ATATATAAAT 60
NCTGCTTGA CTCTAAA 77

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SEQ ID NO:6381
SEQUENCE LENGTH:32
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07546

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SEQUENCE DESCRIPTION:
GATCTCATTT TACAAGAGAA TCTCTCTGCA AA 32

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EP 0 679 716 A1

5 SEQ ID NO:6382
 SEQUENCE LENGTH:83
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07547
 SEQUENCE DESCRIPTION:
 10 GATCTCTAAG TGTAGCTGTA AATTTTGGGG TTAATTTGGC TTATATTGGN CCTTTTAAAA 60
 GNAATAAAGT TTTTAAATGC AAA 83

15 SEQ ID NO:6383
 SEQUENCE LENGTH:74
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07548
 SEQUENCE DESCRIPTION:
 20 GATCTATAAA GAAACATAAG CTTAAAGTTG TTTATCACTG TGGTGTTAAT AAAACAGTAT 60
 TTTCAAAAAA TAAA 74

25 SEQ ID NO:6384
 SEQUENCE LENGTH:342
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07549
 SEQUENCE DESCRIPTION:
 30 GATCCAGACC TCTCTAATAT GAAGAATAAT GAGCCTTATN ACTATAAGTT TGTNAAATGG 60
 ATGACTAAAC ATAAGAACT ATCAGCCATC CCCGTTTCAG CATTCTGTAA CTCAGAGACT 120
 AAATCACAGT TTGAGAAGTT TGTNCGTTTT TCCTTCATTA AAAAAGACAG CACACTGGAT 180
 GCTGCTGAAG AAATCATCAA GGCATGGNGT GTACAGAAGT CTTGATTGT GCAGAATGGN 240
 TTAATGTTTC TGTTAGATGA CCTAGTATGG NATTGTTNACT TAGTGCTGCC ACCTGCTGGG 300
 35 ATGTTAANNG ANTGTNGGAT GCGTGGCCTG GCGGTGNNNG CN 342

40 SEQ ID NO:6385
 SEQUENCE LENGTH:118
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07550
 SEQUENCE DESCRIPTION:
 45 GATCCTGTGT TTTGAGTCCC AAAGGCCATG CCAAGGATTG GCTTTGGNTG GCTTTAATCA 60
 CCAACCCATC AACATCAAGC CTCNCCAGG CCGGTTCAAA TAAATGTANT TAAATAAA 118

50 SEQ ID NO:6386
 SEQUENCE LENGTH:277
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

55

CLONE:HUMGS07551

SEQUENCE DESCRIPTION:

5 GATCCAACAA ATTAACCATA TAAGCACAGA AAATAGAGAA ACACAGTTAT TGAATCTACT 60
 CTTGTCATTA ACATTTTCAA AAAACAAAAT GCATATTGTA ATATTGGTA CATGACACTT 120
 GCATGTTGNT ATGCCTATAT NCTTGCAAAG NATTCAATGT GTACTTAGCG GCGNTTAAAN 180
 CTATGTCATG TACAACNTT ATAAGCNTTT TTNCNGGGGT CCCATTGGC CTCATCTTT 240
 CAGTAAAGTC TTTGTCAGGN AANNGTTGN TGGTTAN 277

10

SEQ ID NO:6387

SEQUENCE LENGTH:102

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

15

CLONE:HUMGS07552

SEQUENCE DESCRIPTION:

20 GATCCAGGGA TGGGAATGGG AGTGGGTAGG AGGGGTAATG CGGTCATTGG AGTCGGGGGC 60
 TGGAACATTA TGAGTGCTCA ATAAATATAA ACTAATGAGA AA 102

20

SEQ ID NO:6388

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

25

CLONE:HUMGS07553

SEQUENCE DESCRIPTION:

30 GATCCAGAGG TGGGTGCAGC TGAAAGTAAA CAGAATGGAT TGCCAGTNAC ATGTATGCCT 60
 GCGCAGTTC CTTTNNATTT GCAGAAGCTG TGAGTTTGT NCACAATTAG GTTCCTAGGA 120
 GCAAAAAACC TCAAGGATTG ATTTATTGTT TTCAACTCCA AGGCACACTG TTAATAAACG 180
 AGCAGGGTGT TTTCTCTCTT CCTNTCTAAT ATATGGAGTT TCGAAGAATA AAATATGTGN 240
 GCAATATTTA AATTCTCAGG AATTGNTCTA TACTCTTNGG GAATGGANTT CAGTTTTCAN 300
 TCANGNTTTA CATTTNTGGT GGCTTAANAA NCNTTGGN 338

35

SEQ ID NO:6389

SEQUENCE LENGTH:245

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

40

CLONE:HUMGS07554

SEQUENCE DESCRIPTION:

45 GATCAGTCTT TTGAAATTGG GATTCTAAAT TAGAATGAGA GGAGGAGCTG CTAAAGCATA 60
 AAAATGGTAG GGGTGATTTT GGTNTTTTAT CCTAGTGACC AATGTTGACC ATTAATTGTA 120
 CATTATAAAT AATGTTTTCN TCTGGGACTT TTGCACTTA ANGCTGGTAA TTNTNATCAT 180
 AGGGCTTCAA ATACAAAAGA NTTGTATGCG GTNTTCAGAA AATAAACATG CTCTTTGCAT 240
 ATAAA 245

50 SEQ ID NO:6390

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS07555

SEQUENCE DESCRIPTION:

GATCTTNACG CGNCTGGGGG CGAAGATTTC GCCATGGATG AGGATGGGGA CGAGAGCATT 60
 CACAACTGA AAGAAAAAGC GAAGAAACGG AAGGTTGCGG GCTTTGGNTC CGAAGAGGGG 120
 TCCCGAGCGC GGATGNTGAG GNTTATNACA GCNTGGAGCA GGATGGCGAT GAACCCGGAC 180
 CACAACGCTC TTTTGAAGGC TGGATTCTNT TTGTAAGTGG AGTCCATGAG GAAGCCACCG 240
 AAGAAGNCAT ACACGTCAA TTCGCAGANT ATGGGGAAAT TAAAAACATT CATCTCAACC 300
 TNGNCAGGCG TTCCAGGNTA TCTGGAAGGG GGTNTNCTCT NGTCN 345

SEQ ID NO:6391

SEQUENCE LENGTH:343

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07556

SEQUENCE DESCRIPTION:

GATCCCCTGC TGGTCTCTGG CAGTCTCCTT GATTTGGGT ACCATGTATA TTTCCCGCTT 60
 TNACTTTAAC GCTTTCTAGG ATAGGGTAAG CACCCTTAAT TCAGGCACCG TCCATNAGCT 120
 TCCTTTGCAA AGGCTACTTA TGGCCGGTCA CAATCCAGCA CTCAGACAGA GCCAAGGCAA 180
 TATCCTCTTG CCCATGGCTA TNATGTCAGA CAGTGGATGG GCTCCAGCAA CAAGAGTCAA 240
 AATAACTAAA GGCCTTTGCT CTCCTCTGAC ATTGAGGCCT GGGGCTTACA GTTTGGAATA 300
 CAACATGTGA NGGTTTTTGN TNGTTGTTG TATTATTGNA GAN 343

SEQ ID NO:6392

SEQUENCE LENGTH:185

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07557

SEQUENCE DESCRIPTION:

GATCAAAAAT TGTGGCCATC TTTGCAAATN ACTACCTNTA GCCTGTNAAA ATACATTTCA 60
 AAAAAATGTTA TGTGCAATGA AACTAAATT NAAGAGCAGT TACAGTGTGA CTCACTCATG 120
 TTTAAAAAAN ATCGAAGAGC TAAAAAATAC GTCTAATTTA TGTAACCCNT TGGNATGTAT 180
 TNNTN 185

SEQ ID NO:6393

SEQUENCE LENGTH:68

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07558

SEQUENCE DESCRIPTION:

GATCATTTCT TGTAATGTTT TGAGAGTAAT GCATACAGAA ATATAATAAA NTGTGTTGAA 60
 ACTGCAAA 68

SEQ ID NO:6394

SEQUENCE LENGTH:163

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07559

SEQUENCE DESCRIPTION:

GATCCCCATT GTGTATGANC TGAACAAGGA GCTGAAGCCC ACCAAGCCCA TGNAGTTCCT 60
GGGTGATGAG GAAACGGTGC GGAAGGCCAT GGAGGCTGTG GCTGCCCAGG GCAAGGCCAA 120
GTNAGGGGTG GGCTTGGGCA ATAAAGGCAC CTCCCCAAC AAA 163

SEQ ID NO:6395

SEQUENCE LENGTH:275

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07560

SEQUENCE DESCRIPTION:

GATCAACTCC CACTCATCCT GCAGATTTAA GCTTGGATGT CACCTCCCGA GGGACGCTGT 60
GCCTGCCTCC CTCGTCAGAG CAGGTGTTGT CATCTACTCA CAGAATCTGG CTTCTTCCA 120
CAAAGCACCC CCGTGAGCAT CTGTTTAAACA TTGTCTCTCC ACCTAGACTG TAAGGTCCGT 180
GAAAGCAGGG ACCACACCCA TTTGTGCTCA CTTGGGACCT ACAGCCTCTA ATCCAGAGCC 240
TGACACAAAC CTCAATAAAT ATTTAAGGAC TGAAA 275

SEQ ID NO:6396

SEQUENCE LENGTH:131

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07561

SEQUENCE DESCRIPTION:

GATCCGGGAC CTTCANACCC AGGGAAAGGG TGAGGGAGAC TGGGGCCTGG TCTGCTTTC 60
CGGGCCTGAA AGCTTCCCGG AGGTTTGAG GGTGAGNAG GAGGAACGGT GGGGGNNGG 120
AGTCACTGNN N 131

SEQ ID NO:6397

SEQUENCE LENGTH:359

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07562

SEQUENCE DESCRIPTION:

GATCCATTTA AAACAAGCTG ATAGTGTTC GTTAAGCAGT ACATCTTGTG CATGCAAAA 60
TGAATTCACC CCTCCACCT CTTCTTCAA TTAATGGAAA ACTGTTAAGG GAAGCTGATA 120
CAGAGAGACA ACTTGCTCCT TTCCATCAGC TTTATAATAA NCTGTTAAC GTGAGGTTT 180
AGTAGCTCCT TGGTTTTGCC TCTTTAAATT ATGACGTGCA CAAACCTTCT TTTCANTGCA 240
ATGCATCTGG NAGTTTTGAT ACTTGTAAC TTTTTTTTTT TTNGGGGNN ANNTTTTNG 300
GGNCCAGGGG TTTTTTTNG GAACCCCTTG GGCNNTGGCC CNGGGGTACC CGGCNAGNN 359

SEQ ID NO:6398

SEQUENCE LENGTH:61

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07563

SEQUENCE DESCRIPTION:

GATCTCTGAC GATACCTGTA TGTTCCTATT GTGTAAATAA AATTNCTGGT ATGAAATNAA 60
A 61

SEQ ID NO:6399

SEQUENCE LENGTH:367

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07564

SEQUENCE DESCRIPTION:

GATCCTGTGG GAAAGAGAAA GGCATCCCAG AAGTTTGACT TGTTTTNCTT TTAGAATCTG 60
TCTCTGCAGG AAAGAAAAGT CATTTTAATA AAAACCAAAA TAATTTATAA TAATTGTNGC 120
CTTTCCTCC CCATCCTCTC CACTAAGCCA GCAACTAAAT GTATTCAGAA AACAAGTTAA 180
TCAAGGAGGG CAGAAGAATT TTTTAAAC TGTAAGAAAN TGTCCTAAG TTTAGATTAA 240
TTACCTGTCA TTTAGGGCAA AANTCCAGTG TCCCAGCAC ATGGTGACCC AACTNTGTCT 300
CACAGTTGCA AAGTGTGTTG GAAAATAAAG TTGCTTTGTG NACTTNGNGG GNAANNCNNT 360
NANNNNN 367

SEQ ID NO:6400

SEQUENCE LENGTH:379

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07565

SEQUENCE DESCRIPTION:

GATCAGGAAC CAAAAAGAAG AATGTACTTC ATCTGGTTGG GCTGGATTCC CTCTNATAAG 60
CCTTCCCAGT TGAAGGCTAG ATGAGGCTAG GCTCTAGCAA GTTGAAGTCA AACCAGCTCC 120
TTCAAGAAGC TTTGAGCAGA ATGAAGTGGG GAGGACCCAG CTTCCAGCCC AGGAAGCCCA 180
CTGTACCTGG AGCCATCTGG GATAAGACTT TGACCCATGA CTCCCATATC CACAGCCTGT 240
CCATCCTAGC CCATCCCAGT TTATCCTGTA TCATTGAGC TGGGNTTCCA CATCCTCTGA 300
GTTGGTAGTC CCATNTCAAG TCNTTNAATA AANGNCTNTT TGGANTATTT GANTTCACAT 360
TAAAAAANA ANAAAGGCN 379

SEQ ID NO:6401

SEQUENCE LENGTH:225

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07567

SEQUENCE DESCRIPTION:

GATCGGGGTA TGAAGTGTGC ACACGCAGCC CAACAACGGG CAGTGGTCTC TGTGCTCCTA 60
GGCATCCAGC ACAGGTTCTG GCAGGGCACC CCTGCTGGGG TTGGGGGCTG GTCTGTGCAT 120
AATCCTGGAC TGTGATGGGA ACAGCCCAGT GCAGTCTAAA CTTCAATTGT GTTGAAACTA 180
CTTTAATAGA CAAAGTAATA AATCATGTTT ATCTATTGAT TAAA 225

5 SEQ ID NO:6402
 SEQUENCE LENGTH:60
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07568
 SEQUENCE DESCRIPTION:
 GATCTTCAGT TGTATTTTNG TGAATATTTT AATACATCTT TTTCAATTTC TGAACAACAA 60

10 SEQ ID NO:6403
 SEQUENCE LENGTH:176
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS07569
 SEQUENCE DESCRIPTION:
 GATCCGGTGC TACGGGAAAC ATTTTCCTAA GATGCCCATG AGAACAGACC AAGATGTGTA 60
 CAGCACTATG AGCATTAAAA AACCTTCCAG AATCAATAAT CCGTGGCAAC ATATCTCTGT 120
 20 AAAANCAAAC ACTGTAACN CTAAATAAAT GTTAGTCTT CCCTGTAACC TTCAA 176

SEQ ID NO:6404
 SEQUENCE LENGTH:360
 SEQUENCE TYPE:nucleic acid
 25 TOPOLOGY:linear
 CLONE:HUMGS07570
 SEQUENCE DESCRIPTION:
 GATCTTCCTT GCGGCCTCTG CCCTGGCCTG CTCCAGCA CACACTTCTT TGGCCTAAGG 60
 GCTTCTCTCT CAGGACCTCT AATTGACCA CAACCAACCT GGGCTTCAGC CACATCAGTG 120
 30 GGCAGTGGAG CTGGGGTGCA CATGGGGCCT GCTCACCTTG CCCACACATC TCCAGCCAGC 180
 CAGGGCCCTG CCCAGCTTCA ATTTACAGAC CTGACTCTCC TCACCTTCCC CCCTGCTGTC 240
 CAGAGCTGAA CATAGACTTG CACTTGGATG TCACCTGGAG TGTNACATGG GAGTGTATG 300
 GCAGCATCAT ACCAAGGCCT ACTGTTGCAC ATGGGGCCAA AACCAGTAAA CAGCCACCTN 360

35 SEQ ID NO:6405
 SEQUENCE LENGTH:32
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 40 CLONE:HUMGS07571
 SEQUENCE DESCRIPTION:
 GATCTNACTA ATAAATGAAG GGAAAAAGGA AA 32

45 SEQ ID NO:6406
 SEQUENCE LENGTH:219
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 50 CLONE:HUMGS07572

55

SEQUENCE DESCRIPTION:

GATCTGGTCA TTTAAAAGTG TGTGGCACCT CCCCACCACC TCTNTCTCTC GCTCGCTTGC 60
 TCGCTCCTGC TTTNACCATG TNATGTGCCT GCTCCCCCTT TGCCTTCCGC CATGATTGTA 120
 NGCTTCCTGA GGCTTCCCTA GAAGCAGATG TCAGCATTAT GCTTCCTNTA CAGTCTGCAG 180
 AACTGTGAGC CAATNAACT TCTTTCCTTA TAANTTAAA 219

SEQ ID NO:6407

SEQUENCE LENGTH:347

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07573

SEQUENCE DESCRIPTION:

GATCCAGTTA CATGTATTAT NNTTTTCCTA CCTTATGGAC TATTTTGGAG GGATAAGCTA 60
 TTAAGACTAA GACTATGNGT GANAGTNGGG GAAGGAGCAG GAAGGGAGGA ACCTGCACAC 120
 CACATTGGAA CCNGCACACC ACATTAAACAC AAAGGCAATC TTCTGGCTCG GACTGTTCTT 180
 TACTACTGTN CTTAAAGAAA ATGTTCAATC TGCTGCAGCT ANCTAGCCTC CATCTNCTAC 240
 ACCAAATACT ATTCCATGCC ATGGAAGTGC TATGCAATAA CTNTCCCAGG TAGGCACCTT 300
 ATACCCGCTT AAANAGNCTT TAAANATNTG CANTTTTGGT NGGNGTN 347

SEQ ID NO:6408

SEQUENCE LENGTH:208

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07574

SEQUENCE DESCRIPTION:

GATCACTCCT TATTTGCTGT TATTTTNGTN CTCTTAACCA TACATTTCCC AATGTTTAAA 60
 GGGTGGTATT TNGAGAAAGT NAATGGGTTT TGTAAGTGCC AGCNCTTTAT ATAAGCAGGT 120
 AGACTAGCTT TTNCCCAGCA GACGCTTGTC TGTNTAGACT TCCTTAAAAA AATTATCTTC 180
 AAATAAAGCA AAATCTCCAT AAAANGNN 208

SEQ ID NO:6409

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07575

SEQUENCE DESCRIPTION:

GATCCATAGA CTTAGCAAGT CTTGCCTTAT CTATGGAGCT CGATGGTGAG AATTGTNACC 60
 ATTGTCTGAT GTCCATAGTT CCTTCCCCT AGATTGTTT TTTCCACGGA TTGTGTTTCAT 120
 CTGAACCAT TTTATTTTAA TTTACCAAAG TACTGTACTT GGCTATTTGC AGTGTTTTCA 180
 AAACCAAAATG TTTCTNTTTT GGTGTTTTTA ATCTCCGATA CTTGGTGCAA TAGAAGCTGC 240
 AAAGATGTGC CACTTNATCT ATGAAATGGA GTNTTGATA CCAATAAATN CTAGTTTAAA 300
 AACANTTNAN NANNNANN 318

SEQ ID NO:6410

SEQUENCE LENGTH:233

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07576

SEQUENCE DESCRIPTION:

GATCTGTAA GGGGTGGATA ACATAATATG CAGCTTAGGA TGCTATTTTA AGATGTATGA 60
TATTCAGTTC ATTACCTGA TTAATTTGGT TGCAGCACAA CTGTATATAT NGTATAACCG 120
AAATTGATTA TTTNCATTGT CCTTAATGCA GTGATTTATA ATTNGAGCAT GTTTAATAAG 180
TTTNCCTCTC TNGTTAACTA GTCATTTGAC TGGNAANNNA TAANATNCTT TTN 233

SEQ ID NO:6411

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07577

SEQUENCE DESCRIPTION:

GATCCCACGA GGACCACTNT GTAATTGNCA GGGGGCTTCC AGNNAAGACA CTGAACCCTG 60
AAGCCGTGGT CTGGGGCAGG GTGGTGGGAG AGGTGGGCGG GGGNN 105

SEQ ID NO:6412

SEQUENCE LENGTH:101

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07578

SEQUENCE DESCRIPTION:

GATCACCTTC CCACTTCATC CATCCCCACC CTTCCAGCTC CCCATCCATC CTGCTGAAAG 60
CATTTCCACC ACTCAATAAA ACCTCGCATT CATCCTTCAA A 101

SEQ ID NO:6413

SEQUENCE LENGTH:332

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07579

SEQUENCE DESCRIPTION:

GATCAGAAGG AATCCGNCGA GGCAGGGATG GGTGTGCCCA TGTNTGCCCTT GACGGGACTT 60
CATCTTATAG ACTGTTAAAC TGTCACACAC AAACAGGCTT TCCACCCCTG CTCTGNNGNC 120
ACCACGCACA GNTTTCAGT TCTTAGTGTG GCTGTTTAAA GTAGAAAATC TGGGGGCTGG 180
GTGAGGCCAC TCATGCCTGT AACCCAGGGC TTTAGAAGGC TGAGGCTGGG GGGNTTGTTC 240
GAAGGTCAGG AGGTTCAAGG CCCAACCTTG GGAACCATAG GCAACNACCC CCCATNNTNT 300
NCCAAANTNG AAAAANCCAA AAAGGNNAAAC CN 332

SEQ ID NO:6414

SEQUENCE LENGTH:129

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07580

EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCTACCAG CATAAATATC TNCTNATTG TCCCTATGCA TATCANTGA GCTTCATATA 60
CCAGCAATAT ATCTNAAGAG CTATTATATA AAACCCCAA ACTNTTGATT ATTAGCCAGG 120
TAATGTGNN 129

SEQ ID NO:6415

SEQUENCE LENGTH:257

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07581

SEQUENCE DESCRIPTION:

GATCTAGAGG AATTAGTCAT GACATCAAGA AAGGAGTCTA AAGAAGAGAA AGAAAATCAA 60
GATGAAAGAG AAGTAAAGA AGANGAACAA GAAGAAGAAG AAGAAGTCAG GTCAGCAGAA 120
ANTTCCTCAA ANTCTCCAAA GAAAGGTCAC TGATTCCAAG GCANCCTGTG GNAAGANGAN 180
TTCAGATGTC AGTGGGAGCC TCCAGGTGGG NGGNGAGCAT GCCTAANCCC TGGTGGCCAT 240
TGTNCTGTAC TANGGAN 257

SEQ ID NO:6416

SEQUENCE LENGTH:138

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07582

SEQUENCE DESCRIPTION:

GATCCACAAG GCTTCTGTAT ACAGTTCACG GTCCTCGGCA TTGCTTCTNG TAATTTNCC 60
ATCCCAAAGG AGAGCTACTG TACTGACTGT ACTTGTGGAA TNCAGCGCTT CATTAAATTA 120
AGTTTATTAA AATGCAAA 138

SEQ ID NO:6417

SEQUENCE LENGTH:331

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07583

SEQUENCE DESCRIPTION:

GATCTAATGG GACACGGCCA GTGCCTAGGG GTGCCAAGTC CAAGGCCTCC CACTGGGNGT 60
AATCGCTGAG AAGATGCCAA TTTTCATCC ACCGGCTGCA CAGGCACAAA CTCCCCACC 120
CAGGACGGCT GTNATGAGGT GGCCCTCCCT NTNAACCCTG GTCCCTGGAG TCCCCAGCAC 180
CTGGGGCCCT GGTGGGGCTG ATGTCACAGG TGTTTACTGT GCTGCTGCAC TGGTCCTATG 240
NCAGCTTCAC CCCATGTGGG GACCACGGAA GGNACAATNC CTTACCCCG NTGCGGGN 300
CCGTGCGGTN NNCAAGANGG GNCAAGGANG N 331

SEQ ID NO:6418

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07584

SEQUENCE DESCRIPTION:

GATCCAGCTT ATCCTTTAAT TTCAAAGTCC ATTCTTGGGG CTGGTGGGGA GGCAGGAGAA 60
 TACCCCTCCC TAAGCCCTTA GTGTGTGCCG AGCTTGCTTT GTNATGTTGG CAGGGGAGGG 120
 GAGACCTGGG TGGTGACTGA GTTCCCTTTA TCAAACCCCTT CAATGGGCAC AAAATTNAGT 180
 GCTTGATTTT ANGGTTTAT TTTTNNATGA ATGTCCAAAA TCTGTGTTTC CCCNNTGCCC 240
 TCCCAAGANN TGTGTGGGCC AGTTGAAAGT GTCTTGGTTT TGTGATCAT CTTTCCCTN 300
 AATTCTGGA GCANGNCCC TGAGNNCCNT GCCANAN 337

SEQ ID NO:6419

SEQUENCE LENGTH:350

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07585

SEQUENCE DESCRIPTION:

GATCATGGAT GTAGCATACA CCAAAATCCA CCGAGACCTT TGCAAAATNA GGCTACTTGT 60
 TTNATTTTGC CAGATTATCA TGGGAGAGTC TTTAACTAGT NCTGAATATT TGTAAGTCTG 120
 AGTACTTCAT AGAATTATAA AGAAGGTACA GCTGACCCCT TCTCACCATG TNGTAAATAT 180
 TTCCTGAGAC TGGGTGCAGT ACTGAGGGAT AATAATCTCC TGNTTTATCA AATCCTGACT 240
 GTCTAGAGCA ANTTGTACAC TCTCTTTGTG AATGGTCTG TAACGTGTAT GNNCACATTT 300
 NTGGGTGCCT TAGANGTNGT NTTTTNCATG TGTGCTAATN TGCAGTTTCN 350

SEQ ID NO:6420

SEQUENCE LENGTH:362

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07586

SEQUENCE DESCRIPTION:

GATCCAGGGT CTNAACCGCC GTGGGACGAG GAGGGCATCC CGGCCACAGC AGGGGCCGAG 60
 GAAGAGGAGG AGACAGAGGG GAAAGGGGAG CCGGCCTGAC CCACACCCCC GGCCATCGCT 120
 CCCTGGGCCA GGTTTAGAGC AGGGAGTTTG GCTGGTGCTG GGCCTGAGCC AGGGGGCCGG 180
 GACACCCITG TTTCCGGTGG TCTTCCCCTT GTGGGAGCAG GTNGAGGGTG GAGACCTAAA 240
 CTTTGGGGTC CAGCTGCCTT CAGCCCNCT CCNCCAGACT AACAGACCCT TGGAGGCAGG 300
 GGCTGTGGA ATAAATCTTT GCCTGNTGGG NTAAAAAAA ANTATNANNT GTGNNNNGTT 360
 GN 362

SEQ ID NO:6421

SEQUENCE LENGTH:114

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07587

SEQUENCE DESCRIPTION:

GATCTNCTCT CCCACCCACA CTTATCTCCC CCAGGGCCAC TCCAAAGTCT ATACACAGGG 60
 GTGGTCTCTT CAATAAGAA GTGTTGATTA GACCTGAATT TCTCCACCTA TAAA 114

SEQ ID NO:6422

EP 0 679 716 A1

SEQUENCE LENGTH:62
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07588
 SEQUENCE DESCRIPTION:
 GATCTTTGCT ACTGGAGTCC TTAAACAACA CCTATAACGA TAAAAAATTC CTAATTNTTA 60
 AA 62

10 SEQ ID NO:6423
 SEQUENCE LENGTH:96
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07589
 SEQUENCE DESCRIPTION:
 GATCCTTCAT ATTTAATTAA AAGTGTTCAT TCAGGTAAGG TGTATGTTGA AATTTNCCA 60
 ATNATCTTAA TAAACCTGG CAATTAAAA NCCAAA 96

20 SEQ ID NO:6424
 SEQUENCE LENGTH:88
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07590
 SEQUENCE DESCRIPTION:
 GATCTATTTA TNTAAGTATG TAGTAAACAA TATCTNAAAG TGTCCGATTC ACTACTTGTA 60
 AATTAAAAAA GTNATGATTA ATGTGAAA 88

30 SEQ ID NO:6425
 SEQUENCE LENGTH:314
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07591
 SEQUENCE DESCRIPTION:
 GATCAGAAGA AGCTACAGTT TTNTTTTAAA CTTGAAAAAA AATTAAGAAG TATTAAGGAA 60
 AAAGCTTAAT ACTCTGGAAA CCAGACATCT GGTCTTTCTC ATGCGTAGGG AGCAGTGTA 120
 ATGAAAGTGA AGTCAAACGG ACAGTTTTTT NCTGCATTAT GCACAAGGTG TTTTAAATTG 180
 TCAGCTTAAG GTTTAGATGA NGATAATCTT TAACTTTGTT TACAATTTTG NCTATTTNGC 240
 ATTATCCCCC TTTTNCCATA AAANNTAANG TGAATATGTG TGCATATNNN TNTNGAGTCA 300
 NAATTAAACT TATN 314

45 SEQ ID NO:6426
 SEQUENCE LENGTH:91
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07592
 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

GATCCTGTGG TGTGTATAA AAATGAACAG CTAAAAATTT ACCCATGACA AGATTAAAGC 60
AAAAATAAAC ACAAAGTTGT TAGACTTTAA A 91

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SEQ ID NO:6427

SEQUENCE LENGTH:122

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

10

CLONE:HUMGS07593

SEQUENCE DESCRIPTION:

GATCTTGTGC TATACTTTT AAAGCCATAC AAATAAGAGT GCTAAACTNT GGACTTAAAA 60

GTAGGTGTNT AAATATTTT AATCAGTATT ACTTGGAAAA TAAATATAA CAAACCCATA 120

15

AA

122

SEQ ID NO:6428

SEQUENCE LENGTH:293

SEQUENCE TYPE:nucleic acid

20

TOPOLOGY:linear

CLONE:HUMGS07594

SEQUENCE DESCRIPTION:

GATCCTAACC CCTGCAGGGC TCAGGGGTCA GCAGGGACCC ACTGCCCAT CTCCCTCTCC 60

25

CCACCAAGAC AGCCCCAGAA GGAGCAGCCA GCTGGNATGG GAACCAAGG CTGTCCACAT 120

CTGGCTTTTG TGGGACTCAG AAAGGGAAGC AGAACTGAGG GCTGGGATAT TCCTCATGGT 180

GGCAGCGCTC ATAGCGAAAG CCTACTGTAA TATGCACCCA TCTNATCCAC GTAGTAAAGT 240

GAACCTAAAA ATTCATCAA ATGAACAATT AAATAAACAC CTGTGTGTTT AAA 293

30

SEQ ID NO:6429

SEQUENCE LENGTH:209

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35

CLONE:HUMGS07595

SEQUENCE DESCRIPTION:

GATCTGAATT TGTAAGTTAT TAATGTGAAA GGGGGAGCAG GAAGCTCCCC GTTTGCACAA 60

AGGGGGATTG GTTTGCACAA AGTTCACTGT AAATACTGTG AGANTNGANT TCTGAAAAGG 120

TCATATTTAG TGATTTCAA AGCACACATT TAANTGANGA CTGACTAAAT AANAGGTACC 180

40

ACACACTTCT NGGTTGGTTT CATTCCAAA

209

SEQ ID NO:6430

SEQUENCE LENGTH:200

SEQUENCE TYPE:nucleic acid

45

TOPOLOGY:linear

CLONE:HUMGS07596

SEQUENCE DESCRIPTION:

GATCTCCCTC AGTNTGCCCC CAGCCCCCAA ACTCCTCCTG GCTAGACTGT AGGAAGGGAC 60

50

TTTTGTTTGT TTGTTTGTG CAGGAAAAA GAAAGGGAGA GAGAGGAAAA TAGAGGGTTG 120

TCCACTCCTC ACATTCCACG ACCCAGGCCT GCACCCACCC CNCACTCCC AGCCCCGGAA 180

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EP 0 679 716 A1

TAAAACCATT TTCCTGCAAA

200

SEQ ID NO:6431

SEQUENCE LENGTH:131

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07597

SEQUENCE DESCRIPTION:

GATCTNATTC CAGGATTGCT CAGAAATTGG AAAAAAACTC ACCTGTGCTT CCCAAATTTA 60
TTTCCAAGA CAAAGATGTC TATAGCACAG AAAAAAATGC ATGANTTAAA AAATCAAGAT 120
GTATTATTAA A 131

SEQ ID NO:6432

SEQUENCE LENGTH:42

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07598

SEQUENCE DESCRIPTION:

GATCCTGAGC TGTCCCATTA AACATGGCCC TGTCTCGGTA AA 42

SEQ ID NO:6433

SEQUENCE LENGTH:332

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07599

SEQUENCE DESCRIPTION:

GATCCACTAG ACAGTTTCA GTTTGCTTGG AGGTANCTGG GTAATCAAAA ATGTTTAGTC 60
ATTGATTCAA TGTGAACGAT TACGGTCTTT ATGACCAAGA GTCTGAAAAT CTTTTGTNA 120
TGCTGTTTAG TATTCGTTTG ATATTGTNAC TTTTCACCTG TTGAGCCCAA ATNCAGGATT 180
GGTTCAGTGG CAGCAATGAA GTTGCCATTT AAATTTGTTC ATAGCCTACA TCACCAAGGT 240
CTCTGTGTCA AACCTGTGGC CACTCTATAT GCACTTNGGT TTAATCCTTT ATTNCAAATA 300
ANTATACTAA AAGNCTTTAA ANNANGNNA NN 332

SEQ ID NO:6434

SEQUENCE LENGTH:224

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07600

SEQUENCE DESCRIPTION:

GATCAATACG GGTCTATTTT NATGTCAACT GAACACTGTA GGGTACCTTC CAGTCTTTTT 60
CAAGATTGTN AAATTGAGAC AAGTAATTGA ATAATTTGTC CTATTTTAAT TTTAAAAAAA 120
GTGAATGGNC TGAAATGTTA AATGTGAATG TNCATTTCTN AATTGCAACT TTNCTACTGA 180
GTGTTTGACAC TATACTTNCT GGAATCTNAT TTAACAAANN TANN 224

SEQ ID NO:6435

SEQUENCE LENGTH:188

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07601

SEQUENCE DESCRIPTION:

GATCAAAAAG CACAAAGAGC TGGGGCAGAG GCAGGAAGCA GGGGCCCTCC TGGCAGCTCC 60
TCTNAGTGGG GAGAGGTTGG GCAGTGAGTG AGGGACCCCT AATGCAGGGN CTAGAAGCCT 120
CANTTTCCC NTTTTACCCT TCCACACAAT AGCCTCTGTA GGNNAGGCTG CCCCATNCCA 180
CCNGGNTN 188

SEQ ID NO:6436

SEQUENCE LENGTH:256

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07602

SEQUENCE DESCRIPTION:

GATCTAGGAC TAANACTTAA AATGTGGTTG GAATCTCACT CAAGGCCTAC CCACCTAAAA 60
CTGCGTTCTA GCAATTGCAT GCTGACTGCA CCTATGAGTC GTAAGNCTGT AGTTGCAAAG 120
TGTGTCTAGT TTNANATTCT TTCTAAATAC TGGTACGGGN AACAGGCATT CCAGNACTTG 180
TGGCTTGCTG GNGGTGGTNT GAGTNATGNG GAATCTTGGG AAGGGGTTCC TGCCCTGGG 240
ANTTGGGGGG TTTTNN 256

SEQ ID NO:6437

SEQUENCE LENGTH:240

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07603

SEQUENCE DESCRIPTION:

GATCCAGGGC CTTTNAACA TCCCCAAAGT CATGGCCATA CTCACCATTA GCCAGTTTCT 60
AACATCTGTT TCAGGGTATC CAGCTGTAGA TGTCTTATC CCCCATACTT GTGAGTTCTT 120
GGGGTTGCTC ACAAATACTA GGGGTTTTTG TTGTATTTT AACAAATA TAATAATGTC 180
ATATTNATNC TCTTTGTAA CTGCTGGTCT TTACANTAAA GGNGTCATCT GCCTTTCAAA 240

SEQ ID NO:6438

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07604

SEQUENCE DESCRIPTION:

GATCAGCACC CTTCTCCACA TCCACATGAC TGGTTTTTAA TGTAGCACTG TGGTATACAT 60
GCAAACATCC GTTCAAAATC TGAGTCGGAG CTAATAATAA AAAATGAAAA ANCAGAAATA 120
AGAAA 125

SEQ ID NO:6439

SEQUENCE LENGTH:321

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07605

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SEQUENCE DESCRIPTION:

10

GATCCGCCTG CNTCGGCCTN CCAGAGTGCT GGGATAAAAG GTGTGAGCCA CTGTGCCTGG 60
 CCACTCCCTT GAGACCCAGT TTCCACGTCT GTNAAATGGG GGAATACCCA CAGTGAGAAT 120
 TAGATGAGAT AGATGCATTT NAAACCACAA ATCGGGGTGC CCACCTGTGG TGGGTAGTTA 180
 GGGTAGCTGT TACAAAGTGC CCCAGCAGAG ACCCCCTTGA CTAACCTTNA TTCTTCCATC 240
 TTTTCCACA GGTGGCAGCT GCCAACAAGA AGCATTAGAA CAAACCATGC TGGGTAAATA 300
 AATTGCCTCA TTCGGTANTG N 321

15

SEQ ID NO:6440

SEQUENCE LENGTH:123

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07606

20

SEQUENCE DESCRIPTION:

GATCTTTGTC CAGAAAATAA AGCTTTCTGT TGATTTTTTC TTTAAGCAAG AGGTAAGGAA 60
 ATATGCACTT AAGANTATGT GTACAAGAAA ATAGTTTCTA AAATAAATTG AAATGTCTGC 120
 AAA 123

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SEQ ID NO:6441

SEQUENCE LENGTH:284

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07607

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SEQUENCE DESCRIPTION:

35

GATCCACTCT GGACACTCAA CAAGACCTGA ACAGGATTTN CCTACCTGGT CCTTACACTA 60
 CATCATCATC ATCTCATGCC CACCTGCCCC CACCCAGCAG AGCTTCTCAG TGGGCACAGT 120
 CTCTTACTCC CATTNTGCT GCCTTTGGNC CTGCCTGGCC CAGCCTGCAC CCCTGTGGGG 180
 TGGAATGTA CTGCAGGCTC TGGGTCAGGT TCTGCTCCTT TATGGGACCC GACATTTTTN 240
 AGCTCTTTGC TATTGAAATA ATAAACCACC CTGTTCTGTG GAAA 284

40

SEQ ID NO:6442

SEQUENCE LENGTH:215

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07608

SEQUENCE DESCRIPTION:

45

GATCTGGTAC GNGGTGGACG TCATGGCCCG ANATGCCAC GGNAACACAG CGCTGACCTA 60
 CGACCGGCAG GCCTCCAGCC ATGGAGTGCA TCAACGTGCT TCTNCAGTAC GGCTGCCCCG 120
 ACGNGTGTGT ATAGTATCTG TTTTATTGA CTGCAGTCTC CTTGGTGCAA AANCAAANTG 180
 GGGAAANTAA GGATAACTCA GNATTTCAAA AGGAN 215

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SEQ ID NO:6443

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EP 0 679 716 A1

SEQUENCE LENGTH:61
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
5 CLONE:HUMGS07609
SEQUENCE DESCRIPTION:
GATCTNAGAC TTCTCAGTC CAGAACTGTA AGCAATAAAT GTCTGTTGTT TATAAATTAA 60
A 61

10 SEQ ID NO:6444
SEQUENCE LENGTH:75
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
15 CLONE:HUMGS07610
SEQUENCE DESCRIPTION:
GATCCTTGCT CTACCTGGCT AGAGTTTCTN CTCATCAGAG CACTGGGACA TTAAACCAAC 60
CTTTTACAAC ACAA 75

20 SEQ ID NO:6445
SEQUENCE LENGTH:207
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS07611
SEQUENCE DESCRIPTION:
GATCTCAGCT CAGAGAGAGA GCATGAGGTC TTTTAACT GTCAGGAAAC AGAGCTGTGC 60
CCAATTCCAC TCAACTTTTG GCACAACTGT TAATCTGGGC CTTACCTAC CTTAACTGA 120
GTTTCTGCAA GCATAGCATT TTAGACACCC TGGATAACC TTTGGGAAT GATGCCACAG 180
30 AATAAAGTTC ACTCTTAACT TTTCAAA 207

SEQ ID NO:6446
SEQUENCE LENGTH:307
35 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07612
SEQUENCE DESCRIPTION:
40 GATCTGATAG TGGTAGGTCT ATCAGGAATA TCAGGNAAAA TNCTTTTAAAT GTTGTGGCA 60
AAGTCAATTC CTATCATTTG TACTTGACAC CAGCTACCTA GCAGGCTCAT TGTGAACTTC 120
AGGTTTCGCC CACGTTATTT ATATAAAGTA TCTCAGTGT ATATAGGGAG TTGCTGGTTA 180
TTTGAAAATA GTAACAAAGG TAACCCAAAC GAGAACTCC ATTTCCATAA TTNGCTTTCC 240
TGAAAGTCAG GCATGTCTTT GAGGACAGGT TTTGGCAGTA TTNNNTNGT GCACAGGNGG 300
45 GAAATTN 307

SEQ ID NO:6447
SEQUENCE LENGTH:219
50 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

55

EP 0 679 716 A1

CLONE:HUMGS07613

SEQUENCE DESCRIPTION:

5 GATCTGCCTG TTTATTTTGG TGGGTGGTCT TTCCTCCCTC CCCTACCACC CATGCCCCC 60
TTCTNAGTCT GCCCTGGCC TCCAGCCCCT AGGGGACTAG CTGGGTGGG GTTCCTCGGG 120
CCTTTTCTCT CCTNCCTCTT TTCTTTCTGT TGATTGTCGC TCCAGCTGGC TGTATTGCTT 180
TTNAATATTG CACCGAAGTT TTTNAAATAA AATTTTAAA 219

SEQ ID NO:6448

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SEQUENCE LENGTH:272

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07614

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SEQUENCE DESCRIPTION:

GATCTGGTGT AGTATATTTT ATCGCATTTT CTTATATTAA AAAATGTCTG CATGATTACA 60
TTTTATTTC TTTGTAATTT ACATTCAGA ATAGTGATT GCTATATGGG TGCCAAGATT 120
GANTATGAAG AACCCGAGTN TTTGTAGTAT TATAGTTTAA AGCAAATCTG TGTGGTGATA 180
CAGCCATAAG ANTGGGGCTT ATATAANCTC TGTACATGTA AGATTTTGTA CAGAGAATTT 240
20 TTAACTTTAT AAATTGTATA TGANCATGTA NN 272

SEQ ID NO:6449

SEQUENCE LENGTH:300

25

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07615

SEQUENCE DESCRIPTION:

30 GATCCACCCG CCTCAGCCTC TGAAATTGTT GGGATTACGG GCGTGACACN CGCGCCTGGC 60
ATGTATATCA TTTTNNATTG TGTATTTTGT AATTTTGTG TGTNAATTTA TTTTAAATAT 120
AGTTTTCAAT TAACTTGTG AGTTTCTGTA GTGGCAAATC CTTGTGTTG CAATATTGGA 180
TTATTTCTCC ATAAGAGTTT CACATTTCTC CTGCTAGGAA ATCCATGGGG AATATTGGAA 240
TTGGGCTAGT TTTAATNNA TNNCTAATGT TTAANGGTTT TGA AAAANAT TGGCGNCAAA 300

35

SEQ ID NO:6450

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

40

CLONE:HUMGS07616

SEQUENCE DESCRIPTION:

45 GATCTCCACT TTNCCAAAGG GCTGCNAGAG GTGATTGANA AGAACCGGCC TGCTGATGTN 60
AGGGCGTCCA CTGCCCCAC ACCGTCCACA GCAGCTGTCT GAGCCCTCAA TCCCCAAGCT 120
GGCAGCTGTC ATTCAGGACC CCAACCCCTC TNAGCCCTC TTTTCCACA TTCATAGCCT 180
GTAGTGCCCC CTCTAACCCC CAGTGCCACA GAGAAGACGG GATTGAAGC TGTACCCAAT 240
TTAATTCCAT AATCAATCTA TCAATTACAG TCCGTCCACC ACCAAA 286

SEQ ID NO:6451

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SEQUENCE LENGTH:97

55

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07617

SEQUENCE DESCRIPTION:

GATCTGTGCA GAGAGAGATG GCAGCCACTC CCTTCAGTCC TCCCAGATTT CTGTAGCTAT 60
TTATGTAGCA GGCTCAATAA AATGTCTTCT CTCTAAA 97

SEQ ID NO:6452

SEQUENCE LENGTH:113

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07619

SEQUENCE DESCRIPTION:

GATCENAGGC ATNNTTCTT CAGTCAGAGC ATTCAATCCT CAAGCCATTC ACACATTTAG 60
GCCTCATGCT TTNTCCTGTC TTAAATCACA TCCATTCCT ACATCATTG AGN 113

SEQ ID NO:6453

SEQUENCE LENGTH:118

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07620

SEQUENCE DESCRIPTION:

GATCATAAGT CATCCCTC CCCTCCAGG CCTCCTGCTA TATTGGGGG ACCTGACTNG 60
NTTNGCTGGA GTCCCATGAG GATGTGGGCC CNTTAATAAN GGATAGCAA CAGGCAA 118

SEQ ID NO:6454

SEQUENCE LENGTH:233

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07621

SEQUENCE DESCRIPTION:

GATCCCAAAA TCCAAGGATG CTCGAATCC CTTATATAAA ATGGCATGGT ATTTGCATAT 60
AACCTACATG CATCTCAAA TATGTTAAAC CATCTCTAGA TTAATTATAA TACCAAATAC 120
AATGTAATGC TATGTAACA GTTGTACAA TGGATTGTCA AAGGCAAGCT CTNCTTGCCC 180
ANTNCGGGCC AANTGNNGG TNCATGAGCA AATTAANTNN NTGCTGGTAT TTN 233

SEQ ID NO:6455

SEQUENCE LENGTH:253

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07622

SEQUENCE DESCRIPTION:

GATCCGAATT TCCTGCTCTA AACCTCCAC TGCTTCTGC TGTTCCTACT GAGGAAAGAG 60
GATGTGTGTC ATTAAGCTT TGTGGGACC TGTACAGTC AAGGTGGGTA TAATTATGTT 120
TAATTTATTG AAGTGTCTT GGGCATCGTC TTTTCAAAT TCATATATNC TATCCAGAAG 180

EP 0 679 716 A1

ANGTGNATTC ATNNNCATT ATNAAAAAN ATAAATATT ACCAGATAAA GCTCANCAT 240
NCNCTCTNCC CCN 253

5 SEQ ID NO:6456
SEQUENCE LENGTH:44
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07623
10 SEQUENCE DESCRIPTION:
GATCTTTCCT ATCAGTGTA TAAAACTCA TATTATGGCA GAAA 44

15 SEQ ID NO:6457
SEQUENCE LENGTH:320
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07624
SEQUENCE DESCRIPTION:
20 GATCAGGCAG GGTGCCATTN ATTGTCTTTC TCTCCTAGCC CCCTCAGGAA AGAAGGACTA 60
TATTGTACT GTACCCTAGG GGTCTGGAA GGGAAAACAT GGAATCAGGA TTCTATAGAC 120
TGATAGGCC TATCCACAAG GGCCATGACT GGGAAAAGGT ATGGNAGCAG AAGGAGAATT 180
GGGATTTNAG GGTGCAGCTA CGCTCACCT AAACCTTTGG TGGCCTGGG TATGTCTGA 240
25 GGCCAGACT GTTAAGCAGG CTCTGCTGGC CTGTTTACTC GTCACCACCT CTGCACCTGC 300
TGTCTTGAGA CTCCATCCGN 320

30 SEQ ID NO:6458
SEQUENCE LENGTH:179
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07625
SEQUENCE DESCRIPTION:
35 GATCCAAATN AGAACAGATT GGTATTGCA GGTATCACAG CCTAAAGAAA TTATCTTTT 60
GCAAAAGAAA TATTAATGA TTTAGCAGTC TCCACGTGTG TTAATGTTT AAACGTGTAT 120
CATAATGTGT ATAATTGTGT AACAAAATTG TCTACAATAA ATCTNTTGGT ATTTGTAAA 179

40 SEQ ID NO:6459
SEQUENCE LENGTH:337
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07626
SEQUENCE DESCRIPTION:
45 GATCCATTCC CCCCAAGGGG ACAACGGGGT GTNAGNGGAG TAAGGGGTGG AGGTAGGGGC 60
CAGAGGGGCT TACACGATAT CCCATACCTT TAATGCCTTT GGCCTTCCAT TCTNATTTCN 120
CTNATGAGAA TATTGCTGGC CTGCTTTCC CTGGTAGGTA TTTGCCAGGC CCAATGCTTT 180
AACCTTAAGC TGATACTTTG CTTTAGATGT CAGTCTCGTT ACCAGCAGCC TTTTGACCCA 240
50 ACTACGGCGC TCTATATTTA AGTNGAGGGA TTTTNANCCA TTGTGCATGG NAAAAAGNTG 300

55

TTCATGGGTA CNATNGTAAG NAAANNTAAA ANTTTTN

337

SEQ ID NO:6460

SEQUENCE LENGTH:101

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07627

SEQUENCE DESCRIPTION:

GATCGGTTGN GNCATGGAGT GGGGGCTTCC AGGTGGACCT NCCGGGCACA CATTCCATTT 60
NTTGAGCCCC AGTCCTGCCC CCCACCCNAC CCTCCCTACC N 101

SEQ ID NO:6461

SEQUENCE LENGTH:138

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07628

SEQUENCE DESCRIPTION:

GATCCTTCTT GGTTAGTATT ACTTGTNCCG GAGCAATTCT NATTCCTCGT TTAGACATTG 60
NNATTTCCTT CGTTGGAGCT GTGAGCAGCA GCACATTGGC CCTAATCCTG CCACCTTTGG 120
TNGAAATNCT TACATTTN 138

SEQ ID NO:6462

SEQUENCE LENGTH:172

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07629

SEQUENCE DESCRIPTION:

GATCTNATTA GNNTAGTAAT GTTTTCCCTT TAAACAGATT CATCATTATC CCTAAAGTAT 60
TGCTGTATTA ATACTGTCTT CTAGAAAGTA TACACCAGTN CCTACTTTCC TCCGATATCA 120
TTAGCTGTTT TCNGAAACTN AATTTCCTCT TCAGAGATTN CTCATATGTN TN 172

SEQ ID NO:6463

SEQUENCE LENGTH:239

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07630

SEQUENCE DESCRIPTION:

GATCCACAGA GTTCATAGGC ATGATTAAAT GGTAGTAGTT TTCCACCACG AAGTTTTGGA 60
ATAATTTGTC ACAAAGCAAT ANGGAATTA GAAAAATAAC CTTTCAGAAG GAGACAGTTT 120
CTCCAAAGCT TGGGTACTTT CGCTTAATAT TAAANAGTGG ANGGAATTTT TTNCCTGCCA 180
GNCAAAAGGC AGNAGGTGTA TTTCTACACC ATGGNTAGGC AANTNNTTTT TTNCCTAN 239

SEQ ID NO:6464

SEQUENCE LENGTH:213

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07631

SEQUENCE DESCRIPTION:

5 GATCTGCATT TAGCAATGTG ATGTCAGTAA ATGGACATAA CAGGATTGTT GTAAAGGTTG 60
 GGCATGATGT ATGCAAAGTA CTGGCCAGGG TAGACTAATA ACTGATGGCA TTTATATGCT 120
 GTGCTGGAAT ATTGTTACCA AGCTGATGTG CCGTTCTCAC CCNGGGNGAA TACTGGTTTT 180
 GTCATTTTCAT AAATGATATT TTTATAANTA AAA 213

10 SEQ ID NO:6465

SEQUENCE LENGTH:329

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

15 CLONE:HUMGS07632

SEQUENCE DESCRIPTION:

20 GATCTGGCCA AGAATGAGGC CCTCCCAACA CTTTCACTCC CTCTCCAAGC CTTGATGGGA 60
 CCTCCACTTA TTTAGGCCTN ATGTGCTTTG AAGAAGCTTT GAGAGCCAAT NTGTCTTCCA 120
 CGGGTCTCTT TTTTGCTACA AGTAATCAGC CCCATGTNTT CTCTTAAACT GAGAATTGCA 180
 CCGGGGGCAA TTCTGTTTT CTAAGGTGGT CTCTGCTGCT ATTTAACAAC CCAGAGTAGG 240
 CCTCTGTNAG GCTTCAGTGG CCTCAGAAAC CAGAGGGTCC AGATAGGGGG CCTGCTTGGG 300
 CCCTCTGCTG NCAAACTGCT TAAAANTTN 329

25 SEQ ID NO:6466

SEQUENCE LENGTH:236

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

30 CLONE:HUMGS07633

SEQUENCE DESCRIPTION:

35 GATCCACGTG TGTGGCCATA TTGTAACACA TTTTCTGCA AATNACCTCT TTCATTTAAC 60
 AGCCCTTATT CAATGGCCTT TTNCTTTTTC AGTAGTACAT ACACATCTGT GTCATTNGTT 120
 GAATGACGNC ATGAATGTTT TGTAGATAAA ATATAATTAA ANGNTTTTAA GCATTTCTNA 180
 CTCATAGATA TTNATGTTAG CACCAAATTT GNTGTGCAAT GANCCTCCCT ANATAN 236

SEQ ID NO:6467

SEQUENCE LENGTH:239

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS07634

SEQUENCE DESCRIPTION:

45 GATCTATGCC TTAGTGTGTG TTTTNTTGTG TGCTTTGATG TAAAAGCAAG AGTATTTGGA 60
 GCAAGAAGCT GCAATGCGCC AAATACACAN TAGGCCTTTC NTTTATCATT CATACGACAT 120
 TTCTGGTTTA GGTGTATGTA GTTGGGCCAT GTTACTTGTC CAGAGGAAGA GNCTAAAGAT 180
 ATACGGNTAA TTTTNTAGTC CNTAATGCTG TTTTGNTTAC NTNCTNTCAN GGTTATTTN 239

50 SEQ ID NO:6468

SEQUENCE LENGTH:82

55

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07635
 SEQUENCE DESCRIPTION:
 GATCGAGCAA TTGCAATCCA GCCTGGGCAA CAGAGCGAGA CTCCATTTC AAAAAAGAAT 60
 CAATAAAAAAT GTTAAAAGCA AA 82

SEQ ID NO:6469
 SEQUENCE LENGTH:348
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07636
 SEQUENCE DESCRIPTION:
 GATCCGTAAG GCCATGACCA AGGAGGCCAT CCGAGAGCAC CAGATGGCCC GCACTGGCGG 60
 CACGAGACAG ACCTGTTTAC CTGCGGCAAG TGCAGGAAAA AGAACTGCAC CTACACACAG 120
 GTGCAGACCC GCAGCTCTGA TGAGCCCATG ACCACCTTTN TTGTCTGCAA CGAGTGTGGA 180
 AACCGCTGGA AGTTCTGCTG ACCCCTTGTG TAGATGTGCT GCAGCCTTNG GCCTTCNCGG 240
 GCCACGTTCT TCGTTGACAN AGGTTTNTTT GGGGNCCCCT AGGAAGGCGG TATGTNCCTN 300
 GCCNTTNAAN CTGNCTGGN TTGGATTGNA NCTTTTTTGN CCTTTTTN 348

SEQ ID NO:6470
 SEQUENCE LENGTH:178
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07637
 SEQUENCE DESCRIPTION:
 GATCGTGAAA TCTAAGGCTG CTTGATGTTT TGATGCTTGT NAATGCCGTC GTGCTCTGTG 60
 ACTCCACGAA ACGCCAAGGA GCTTCTTCCG GACGNCTCCT CTCCTCTGT AACATACGTG 120
 TGAATAGTCA AGGTTTAATT TTNCTTTAAT CTAATAAAGT CGAAGTGGGA CTNTTAAA 178

SEQ ID NO:6471
 SEQUENCE LENGTH:50
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07638
 SEQUENCE DESCRIPTION:
 GATCCAGCAC ACCTCCTCGG TTTTTTTTTC AATAAAAGTT TAGAAAGAAA 50

SEQ ID NO:6472
 SEQUENCE LENGTH:108
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07639
 SEQUENCE DESCRIPTION:
 GATCTCTGTG AAACAAAATN AAGCAAGAAC GGAGACGCTG AATAGTTTTA TTTCTNTATT 60

AGAAACTGTG ATTGGAACAA TTGAAGAAAT AAAATCATAA CAGAGAAA

108

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SEQ ID NO:6473

SEQUENCE LENGTH:210

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07640

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SEQUENCE DESCRIPTION:

GATCTGAAGA AAGCAGCAAT ATCTGTNACT AGAGAACATT CCCATGTTTT TAAACTCTTC 60

ANTTCTTAGA TGCATTTAAN TTCTTAATGC AAATNACGTA GCAATTTGAA AACTTCTCCG 120

TATTACTNGT GTTTAAATG TCTTGCTTTA AATACAANNC AAATGGTAAA GGGGATTATC 180

15

TTTGGTTTAG ATGGTTAANT ATNATTTNGN 210

SEQ ID NO:6474

SEQUENCE LENGTH:307

SEQUENCE TYPE:nucleic acid

20

TOPOLOGY:linear

CLONE:HUMGS07641

SEQUENCE DESCRIPTION:

GATCAGCAGG ACTGTCTCCA GCACGGAGCT GACACTGTTC AGCTGCCTCA ACTGGTAGAC 60

25

GCTCCCAAGA AGTCAGAGGC AGCNGTCGGG GCGGAGGTGT CCATNACCTC CCCAGGACAG 120

AGCAAAAACT TCTCCCTCAA GAACAGAAAC GTTTACCNN CTATTGTATG ACCTTTGCTG 180

AGGGTATGTC CTGCTCCTTT CCACCAGTGA TTTGTATTAA GNCAGCACTT ATATTGTACA 240

ATACTTCAGN CTGTTTTTTT TAAATACATA AAACCTTATG TAAAAAACT CTATTANCA 300

GGGCAAA 307

30

SEQ ID NO:6475

SEQUENCE LENGTH:140

SEQUENCE TYPE:nucleic acid

35

TOPOLOGY:linear

CLONE:HUMGS07642

SEQUENCE DESCRIPTION:

GATCACAGGT TAGGGGTTTC TCTAGATGGG GGTCTGAAA TTTGCAGTGT CTGCTCCTGG 60

40

GAGGCAGCAC CAGAAAGGGC ACTGAAATGT ACTAGCTGGA TGTNACCCAG TCTTAATAAN 120

CAGGTTTTCT AATCCAGAAA 140

SEQ ID NO:6476

SEQUENCE LENGTH:187

SEQUENCE TYPE:nucleic acid

45

TOPOLOGY:linear

CLONE:HUMGS07643

SEQUENCE DESCRIPTION:

GATCAACTTT GTATATTCAC GTGTATTTAA ATATTGTGCA CTAAATGTTT TGCCCTTGTT 60

50

TGCTATTATA TGGTCAAGGC ATTTATCAGC ACTATTGTAA TGAACATG TAAGTGGCAT 120

GGGTCAGGGA AAATTATTC CTACTTTNCT GCCTAATTAA ATTNCTGTTT TCCAGTATTA 180

55

CATTAAA

187

5 SEQ ID NO:6477
 SEQUENCE LENGTH:113
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07644
 10 SEQUENCE DESCRIPTION:
 GATCTTGAT ATTTNATTTT TAAAAAACTA GAATAAACAG AGAGGCATAA NCATATCTTA 60
 GAGTCCAAGT GGTAGTGTTC AGCATTGGAT ATAATAAATG GATGTTTTAC AAA 113

15 SEQ ID NO:6478
 SEQUENCE LENGTH:276
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07645
 20 SEQUENCE DESCRIPTION:
 GATCTCATT TGGGCTAAAA TAAGACAATA TTCAAAGGTC AGAGATATCT AGCCAGAATC 60
 TNATGGAGGC TGGATTTCAG ATTTTNTNAC AGAATTAGAC AGAGGAACAC AGAGGGGACA 120
 GGCTCAGTTA GGGTGGAGGT GTGGGGTAGG GAAGCAGGNC TTGATATAAN TNATTGGAAT 180
 CATTGTCTTT TAANCCAGTG GTTATGTCA GGGTATAGCG TTTCANGGGA TTTGAGGGTC 240
 25 AGATGGGGNA ATGTAGCCTC TTATTTTGCC AGTGTN 276

SEQ ID NO:6479
 SEQUENCE LENGTH:160
 30 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07646
 SEQUENCE DESCRIPTION:
 GATCTGTAC TGTNTCCCGT CACGCCTTGA TGGGACCGTC TAGTTGCTGG AAAACAACT 60
 35 CAGGGCTCCC ACTGATTCTA CATTATGGTG TGTGTACAA TTATTTTCATT ATATATNACA 120
 AGGTAATANT AATAGANATA AAATCCACAG TAANTGCAAA 160

40 SEQ ID NO:6480
 SEQUENCE LENGTH:158
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07647
 SEQUENCE DESCRIPTION:
 45 GATCATAAGT NTTAGGACCA AAGGTGTTTT TAATGAGATA ATCTTTTNA TTTNNTAACA 60
 AGAAAGGAAA TATTCATTTT ACATTTTAAC ATCAACTTTT AAGTGATATT CAGANTTAAT 120
 TGACCATGGT GATACGTTGT AAAANGTTN CATATTAN 158

50 SEQ ID NO:6481
 SEQUENCE LENGTH:112

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EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07648
 SEQUENCE DESCRIPTION:
 GATCCAAAAA AAAAAAAAAAC GAANANAATT NATCCTCACA AATTGGTGTT CTAAATATCT 60
 TAAGANCCTA ATTAAATAGC TGA CTACAAA AAAAAAAAAA AAAATNCAAA AN 112

SEQ ID NO:6482
 SEQUENCE LENGTH:105
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07649
 SEQUENCE DESCRIPTION:
 GATCCTGTGT GGCCGGTGTT TGCATCTTCA GTGGCCACAA GCATAATAAA GCCCCTTTGC 60
 CTTTCTCTGT ATTATATTCA ATACAATACA TCAATAGTCT TGAAA 105

SEQ ID NO:6483
 SEQUENCE LENGTH:252
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07650
 SEQUENCE DESCRIPTION:
 GATCAGATTT AAAGNCTGGA TTCTATCTAC ATAAGTCCTT TNAATTCCAC CAGGGCCAGA 60
 GCAGCTCCAC CACTGTGCAC TTAGCCATGA TGGCAACAGA AACCAAGAGA CACAATTACG 120
 CAGGTATTTA GAAGCAGNGG GNCAACCAGA AGGCCCTTAA CTNTCACCAG TGCATCACAT 180
 CTGCACACTC TCTTCTCCAT TCCCTAGCAG GGGNGGCTAG CTCATTTNAC NGGGTNNAGA 240
 ANCTGAGGNC CN 252

SEQ ID NO:6484
 SEQUENCE LENGTH:36
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07651
 SEQUENCE DESCRIPTION:
 GATCACGNGT AAAGAGAACT GTTTCCTGCC CTAAAA 36

SEQ ID NO:6485
 SEQUENCE LENGTH:125
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07652
 SEQUENCE DESCRIPTION:
 GATCAGCCTC ATCAATATTG CCGTCATCTG TTTAACA CTCCAGTATATT TCCTCAATGT 60
 CTGTTTACTT AAAATTTTGT GGAGTGACAT AATTAATAAG NAATAAAGTC TGAATTATAC 120
 ACAA 125

EP 0 679 716 A1

5 SEQ ID NO:6486
 SEQUENCE LENGTH:128
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07653
 SEQUENCE DESCRIPTION:
 GATCCACACT GGGGNGAAGC CTTATNAGTG TACGANATGT GGGAAAGAGCT TCAGTCAGCA 60
 10 TCGGGGCCTC ANCTCCCACC AGAGACTCCA CACCGGNGAG AAGCCATATA AGTGTAAGGN 120
 GTGTGGGN 128

15 SEQ ID NO:6487
 SEQUENCE LENGTH:329
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07655
 SEQUENCE DESCRIPTION:
 20 GATCCAGAGA GCTTCCTGAA GGACCTGTTG AACTCAGTCC CCTGACCACC ACACAGCAGC 60
 TCGGGCGGCG AAGAGAAGCT GGCTTGCCTT CCACCCTCTG TTCTCCCTCC TTGTGCATTA 120
 AGTTCCCTCC GNGGGATGCT GCATTGGTAC CCAGCCCTCC NCTATCTCAT TTTTCTTGGN 180
 GTGGCTTGGG GTTTTtaggc TTCCNGTTTT ATCTCGTGTG TGTGGTGCAC CAGCTATGAG 240
 GTTGTCTGTA ANCNAAGCCA TCAAAGGGNC TNTAAATANC TAGGNGCCAT GAGTNGTCCC 300
 25 GGNCANCTTT CANAACTTGA GTNTGANAN 329

30 SEQ ID NO:6488
 SEQUENCE LENGTH:101
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07656
 SEQUENCE DESCRIPTION:
 GATCTGTACT GTAATTTTGT TTGTAATCCT GTATATNATG GTGTAATGCA CAATTTAGAA 60
 35 AACATTCAATC CAGTTGCAAT AAAATAGTAT TGAAAGTGAA A 101

40 SEQ ID NO:6489
 SEQUENCE LENGTH:215
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07657
 SEQUENCE DESCRIPTION:
 GATCATGTAT ATAATTGTAA CATCACATTG ATTTTACGGA AGATGTTATA TGGNCTTTAA 60
 45 TGACACAATG TTTAGNGATA AAATGTACAT TATTTTGGTT CAGTTTTTTA AAAAAANTAT 120
 GCTTTAACAA AATTCTTAGG AATNCTTTA AGCAATGCAG GTATTGCGAT ANCTGTAGNT 180
 TTTACAATAA TGTTACTCTA CAGNTGGGAN AATNN 215

50 SEQ ID NO:6490

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SEQUENCE LENGTH:364
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS07658
 SEQUENCE DESCRIPTION:
 GATCTCTTTG TATTCTTTC AAGCCCAAAC CAATTCTGTN CCTTCAATCT AAATAGTGGT 60
 AATATGAATG TTAAAGAAAT GAAATAAGAA ACATGTGCAG GCACTTTGGA AGGTGCTAAG 120
 10 TGA CTGCCCT AAGGAATGAA AAGCAAGGGC CAGGTGGGAG TAGCCAGCG AAGGCACTTG 180
 GGCTGCCAGG AACAGGNGGC GTGGGAACT CTGGCTTAGG NAAACATGAA CACNGGGGCA 240
 ACAGCGGCAN ACTGTTGTTT GAGTTAAATC TAANTCTCAG GCTCTTTNAC GGTAANGGG 300
 15 TTANGGATAC TCCATTNGA TGAAGANAAG NGTGAGGCTG ANNGTAANGC ACATGGCAGN 360
 TNNN 364

SEQ ID NO:6491
 SEQUENCE LENGTH:85
 SEQUENCE TYPE:nucleic acid
 20 TOPOLOGY:linear
 CLONE:HUMGS07659
 SEQUENCE DESCRIPTION:
 GATCTTTCTT CTCCTTTAAG CTCCTTGTTA ACATTTCTCT AGACGCAATT TATTGAAAAT 60
 25 AAAGATTNC TGTTACTCA TGAAT 85

SEQ ID NO:6492
 SEQUENCE LENGTH:122
 SEQUENCE TYPE:nucleic acid
 30 TOPOLOGY:linear
 CLONE:HUMGS07660
 SEQUENCE DESCRIPTION:
 GATCCTCATT TATGTTTGTG GTTGGAAGC AAAGCTAGGT AGCCATTCT TCTGTTCTAC 60
 35 CAAGTTATAA TAGCATTCTAT TTCCCTTTAT ATTTCCCTGA AATAAGCAC ATTCCACTTA 120
 AA 122

SEQ ID NO:6493
 SEQUENCE LENGTH:362
 40 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07661
 SEQUENCE DESCRIPTION:
 GATCCTACAT TAAGGTCATT TGTNAGAATG ATGTTTGTGT TTGNTAGAG TTGGCTGACC 60
 45 TCCAATCNT GGGGTCAAGG AATCCTACTG CCTNAGCTTC CCAAATAGCT AGGACTATAG 120
 GCATGCACCC GGCCATGTGT TTTATTATA GCTCTTAAAG CCCAGATGAA GAAATCACAT 180
 TTTNCCCAT AGTGAAGAAA CATTGGCCA TTGCTTAGTC CTTNNGGGG GTGNETGTCC 240
 TGTTTTTATT AGATTAGNGA GACCCTGTGT GGGCCACAGT TAATATANN CATTATCACT 300
 50 TTAAAGGTAA CTGCACATC TTAGGTTTCA TAATTNCCTT ATTGTTCTNG NCTCAAANTT 360
 GN 362

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EP 0 679 716 A1

SEQ ID NO:6494
 SEQUENCE LENGTH:328
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07663
 SEQUENCE DESCRIPTION:
 GATCGGGGCA CTGCAGGGGC CGAGCCATTT TGGGGGGCCC CCTCCTTGC TCTGCAGGCA 60
 CCTTAGTGGC TTTTTCCTC CTGTGTACAG GGAAGAGAGG GGTACATTTC CCTGTNCTGA 120
 CGGAAGCCAA CTGGCTTTC CCGGACTGCA AGCAGGGGCTC TGCCCANAGG CCTCTCTCTC 180
 CGTCGTGGGA GAGAGACGTG TACATAGTGT AGGTCAGCGT GCTTAGCCTC CTNACCTGAG 240
 GCTCCTGTGC TACTTTGCCT TTTGCANACT TTATTTTCAT AGATTGAGAA GTTTTNTACA 300
 GAGANTTAAA AATGNAATTA TTTNTAAA 328

SEQ ID NO:6495
 SEQUENCE LENGTH:351
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07664
 SEQUENCE DESCRIPTION:
 GATCCTGTTT CTACCAACAC TGCACCTTAT CCCAGGAACC TGCCCTAGAC CTCCAGAGAC 60
 CATATTTNNT CTCCTCCAT TTCTACCCAG ACCTCCAGGC CTCCTTCTGG AATNATAGAA 120
 CCGTAGAATT GGAAGGAATT TNAGAGGGGN TCTAGTTGGA GTTGTGTCCA ACAGAATTCA 180
 TTAACACCAG CCTGGGCTTG TTTTCTCTCC TCCCTCTGGA CTTTTTAAT CTTTCTCTCC 240
 ACCTCAAAAA ATACTTACAC ACAGATTCTT CTTGTACAGG CATCAAAACC AACTNCTNTG 300
 CCCCTAAGGC TGTGTNNGNG TGGTNTCCAG NNANCCNTAC CNCAGTCANT N 351

SEQ ID NO:6496
 SEQUENCE LENGTH:122
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07665
 SEQUENCE DESCRIPTION:
 GATCTGGTAG GGGAGAAATG GCGAATGTAA TACACATGAN ATGGTATATC CTTGCAATGT 60
 AAGGTATCAG AAGGTGGTTT GACAGCATCA TAAACAGGCT GACTGGCAGG AATGAAAACA 120
 AA 122

SEQ ID NO:6497
 SEQUENCE LENGTH:227
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07666
 SEQUENCE DESCRIPTION:
 GATCANTTGC AAACCTCCCC CAGCCCCTAC CATCCAATNT GGACGGAAAA CANGAACTGC 60
 CTGAAGAAGA GTCCAAGCTA CAGATACACA GCGTGTGCAT TGCGGCTGTC ACCTTCCTNC 120

TNCCACTTCT GTATCCTCAG AGATGCTGCG TGGATGTTTC CTTANCTCA GCTGACTTCC 180
CTGTGAATNT CTAATGCTAG TTCANGGGCC TNCNGGAAAT TGANTTN 227

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SEQ ID NO:6498
SEQUENCE LENGTH:238
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS07667

SEQUENCE DESCRIPTION:

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GATCTTGCCC TTCTACTCC AGAGGCTAGT GGTACAGAC AGCTGGGAAT GGCAGCCACA 60
GAGGGTCCCC TCTGGGAGAA ACAGCTTCAC CCCAGCCTCA GGGCCCTGGG CCATCACTGC 120
AGTGGCCCTG GGAGGTGAGG AAGAAGCTGG CTAGAGGAGG GGGCTCCAC CTACCTTTTA 180
TTTAAGCCAG TATTCTTTGT TCCTGCTTGT AATAAACTT CAGTTTATAA GAGTTAAA 238

SEQ ID NO:6499
SEQUENCE LENGTH:142
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

20

CLONE:HUMGS07668

SEQUENCE DESCRIPTION:

25

GATCTGTCAT AACTCATTTT AAAGTGTGAA ATATGCCATA TGAAGAGGGN TTAAGAGGCT 60
GNGAACTCA TATTCAACC AAATCCAGCA TTAGTTTTTC TTAGGTCTAA TANTTCCTNG 120
TTAATAAAGA TTAAATGCA AA 142

SEQ ID NO:6500
SEQUENCE LENGTH:160
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

30

CLONE:HUMGS07669

SEQUENCE DESCRIPTION:

35

GATCTGTCAT TGTCTCCCAT CACCCCCAAA TGGGCCCTC TCGTTGCAGG AAAACAAGCT 60
CAGGGCTCCC ACAGTTTCTA CATTATGGTG AGTTGTATAA TTNTTTCATT ATATATTACA 120
ATGTAATAAT AATAGAAATA ANGACACAA TNNNTGTAAA 160

SEQ ID NO:6501
SEQUENCE LENGTH:266
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

40

CLONE:HUMGS07670

SEQUENCE DESCRIPTION:

45

GATCTGCCCC AGACTCCCAG AGGGTAATGC CACTCCCACA ATCTCAGGCC TGGTACCCAT 60
CCTNTTTCCTA CTGTGAGCAG AGCCAGAAGG TAGGTCTNTT CAGAGTCTGT NCCCTGGAC 120
CTGGGGNGTG GATATGANAT GGGATATCTC CTTCCATTCC CCGNTCCAGG GGAGAGTCAC 180
TAGTTGTACC CTAATCCATT AGGTCCCAA TAGGGGCCCA ATTTACCTG TATNAGGNTC 240
TGTGCATNCC AGCTGCCNNA AATNTN 266

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5 SEQ ID NO:6502
SEQUENCE LENGTH:119
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07671
SEQUENCE DESCRIPTION:
10 GATCCAGACT TAGGGACGAG GCTGTCACGT GTGGGCACCC TCTGTTCCCTG TTTGTGTGTT 60
TGAATAGTCT GAAATNCTGT GACTTTTTTT GTGTGAATAA AGNTATGAAA CTTCTGAAA 119

15 SEQ ID NO:6503
SEQUENCE LENGTH:90
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07672
SEQUENCE DESCRIPTION:
20 GATCTTCGTG TGAGCCGCGN CGTCCGNCGN GACCGAGCTG CCTTCGCGTG CCCCCGNCCT 60
GTGTTTATA AAAAGAAAGA TTCTCGGAAA 90

25 SEQ ID NO:6504
SEQUENCE LENGTH:62
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07673
SEQUENCE DESCRIPTION:
30 GATCTGTCAG TGTATCAAAA CAACATAAAT AAACCTGAAC AACTGATACG CAAAAAACA 60
AA 62

35 SEQ ID NO:6505
SEQUENCE LENGTH:72
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07674
SEQUENCE DESCRIPTION:
40 GATCAAGTTT GGTCAGGTTT TGTACCAAG CTTTGTAAT AAACCTGGTTT TCATAGCTTT 60
TTGGAGATGA AA 72

45 SEQ ID NO:6506
SEQUENCE LENGTH:221
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07675
SEQUENCE DESCRIPTION:
50 GATCTTTTTT CAGAATGTTG GAAAATCCTG TAAATGCAAA TAGTCAATAC TGTATTAAAT 60
ACGTGCACTT GNGTNTGCT TCGCTTGAC AGTTGTAAAT AATCAGAACA TATGNAAAAG 120

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EP 0 679 716 A1

GTACCCTACA GAGGAAATTC TGATACAGAT TATTGCTATA TTATAACCTN TTGCTGTTGA 180
CGGGGCTGTA GGTAANCTAA TGTGTGGTT TGA CTATTAA A 221

5 SEQ ID NO:6507
SEQUENCE LENGTH:42
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07676
10 SEQUENCE DESCRIPTION:
GATCTTCATT TTTGTAATAA AACGGNAGAC TCATCCAGTA AA 42

15 SEQ ID NO:6508
SEQUENCE LENGTH:62
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07677
SEQUENCE DESCRIPTION:
20 GATCTTAGAC TTCCCATCCT CTAGAACTGT GAGAAATAA TTCCTGTTGT TTATGAGCCA 60
AA 62

25 SEQ ID NO:6509
SEQUENCE LENGTH:117
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07678
SEQUENCE DESCRIPTION:
30 GATCTGCCAT CTAGAACTCA TATTCTAAAG GGCAGTGNTT TCTCAGAACA GTGATGTTCT 60
GGAATATGTA TTATTATTT TAACACTTTT TTAATAAANT CTTTATTATA AACCAAA 117

35 SEQ ID NO:6510
SEQUENCE LENGTH:113
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07679
SEQUENCE DESCRIPTION:
40 GATCTGTTTA AAAGATTAAA TCTCAAGTTA ATTAAAAATA AGCTTTTCAA AAATGTATTA 60
TATTTATAAC AAATATACTG TAAATAGANT AAAGACATGC TATTCAGTGT AAA 113

45 SEQ ID NO:6511
SEQUENCE LENGTH:159
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07680
SEQUENCE DESCRIPTION:
50 GATCTGTCAC TGTCTCCCAT CACCCTAAGA TGGGNCCTGC TAGTTGCAGG GAAAGAAGCT 60

55

CAGGGCTCCC ACTGATTCTA CATTATGGTG ANTNGTAGNA TTATTTTATT ATATATTACA 120
 ATGTCAATAA TAGAAATAAN GTGTGCANTA AATGTCAAA 159

5 SEQ ID NO:6512
 SEQUENCE LENGTH:219
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07681
 10 SEQUENCE DESCRIPTION:
 GATCAGCCGG AAGAGAAAAT NAAGACCCCA GAGACATTTA TTGGGGAGTA GGATGTGGCT 60
 GAGTGCTTTT TTTTGGCCA GACTAGCGGA TTCAGTCCTG GAAGAGAGTA TCATATAATG 120
 AGNCCACAG GCACTGGCAC CCTTGGGTTG GNAATAGANG NNGNCCATGG ANTGGCGAAA 180
 15 ACCAGGGATT CCAGCTGGGG NATAGTANCT TGAAGGTGN 219

SEQ ID NO:6513
 SEQUENCE LENGTH:217
 SEQUENCE TYPE:nucleic acid
 20 TOPOLOGY:linear
 CLONE:HUMGS07682
 SEQUENCE DESCRIPTION:
 GATCTTGACC TACAGTTTGA AAAACAGAGT ACTGAAGAGT CTTGTTTCCT TTATTTAGAA 60
 25 CAGTGCAAGG ATGAGTATCT GGGTAGAGTT GTGCGTTGTA CATAACCTTT GATAGTCATA 120
 TAGTTGAACC AAAAGCAANN TCTGTAGTGN CATTGAAAAA ACATTTTGAC TTGATTTATC 180
 CTGNCAGNNT NCNNTAATTT TTGGCTGGCT AATNCAN 217

SEQ ID NO:6514
 30 SEQUENCE LENGTH:119
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07683
 35 SEQUENCE DESCRIPTION:
 GATCTCTGCT CACGTTGCCT CCCTTCATCT TCCTGCATAA GTGGAAGCAG CCTGAAGCCC 60
 TCACTGTTTC TTGTACAGCC TGCAGAGCCG TGAGCCAAAT AAAGCTCTTT TCTTTTAAA 119

SEQ ID NO:6515
 40 SEQUENCE LENGTH:173
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07684
 45 SEQUENCE DESCRIPTION:
 GATCTGATGG TTTTATAAGC ATCTGGCATT TCCCCTGCTT GCACTCACTC TGTCTGGTG 60
 CGCTGTGAAG AAGGTGCCTG CTTCTCCTTT GCCTTCCACC ATGATTGTAA GTTTCCTGAG 120
 GCCTCCCTGG CAATGCAGAA CTGTGAGTCA ATTAACTTC TTTCTTTAT AAA 173

50 SEQ ID NO:6516

55

SEQUENCE LENGTH:238

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07686

SEQUENCE DESCRIPTION:

GATCACCATG TNAGCCTTAG ACCAAGAAGC TGGCTAGTCC TTNCTGTGAA GCTAATACAA 60
 TNGNCATTTC CAGACAAATT TAAAGGNAAC ACTAAGGCTG CTTCAAAGAT TATCTGATTC 120
 CTTTAAAATA TATGTCTATA TACACAGGCA TGCTCTTTT TTAAGTGCTT ACATTTTAAT 180
 AGAGATGANT CAGTTTTNGA ATCTANGCTG TTTGCCCAAG NTGANGCTNC CAGGGTTN 238

SEQ ID NO:6517

SEQUENCE LENGTH:289

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07687

SEQUENCE DESCRIPTION:

GATCTGTTTT TNACCACTTC ACTGAAAGAC ACCATTTATC TACCCAAGGG CAGAAAGTAG 60
 AACTTACTAT TCATTAAATG TTTGACACAA TTGGAATTGT CTTTAATTC TGTCAGAATG 120
 CTATTGAAAA TGTGAATTGC ATGACTTGGT AGCATATTCT TTTCTGCAA AAATAGGCCA 180
 TATTAACCAT GCTTATGACC ANTGNCTGT GCTACTGGCT TTTGGNAAAA ATGGTTTGCN 240
 TCAGNTTGGG GAAGTAATNA AAGGANTNCA CCTGGNGNCC CNNGAATAN 289

SEQ ID NO:6518

SEQUENCE LENGTH:236

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07688

SEQUENCE DESCRIPTION:

GATCCCCAG GATTCTATTN TGTTTAATGG GCTTTTCTAC TAAAAGCATA AAATACTGAG 60
 GCTGATTAG TCAGGGCAAA ACCATNNACT TTACATATTC GTTTTCAATA CTTGCTGTTC 120
 ATGTTACACA AGCTTCTTAC GGTTTTCTTG TAACAATAAA TTTTTGNGT AAATANTGGG 180
 TACATTNTAA CAAACTCNGT AGTNCAACCT AACTNGTAT NAACGGTGTG TAAAN 236

SEQ ID NO:6519

SEQUENCE LENGTH:157

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07689

SEQUENCE DESCRIPTION:

GATCAGGTCT GCTTTNAGTT TCATTTTGT TTCTTTCCCG TCCCACTCTT TAAAACTGG 60
 TTCCGTGAGA AANGGCAGAA GCCGTTCCGT GTCTCTTGCA GGCTGGGCCG NCTTCATGNA 120
 GTNCGAGGCG GTNCCGTGAC NACGTACATA CGTATGN 157

SEQ ID NO:6520

SEQUENCE LENGTH:195

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07690

SEQUENCE DESCRIPTION:

GATCTTTCTG GATATAATGA AGCTTCCTGG AGAAACAGAC TGGCAATCAG AAAGATTGG 60
CTGCTTCCGA CACCTCCATC TTGGGCTGC AGAGGCTGCA GCAATCTGTG AGGCAGCCGA 120
ATCTGCGNGG CAGNCCACTC TTCTAGGACA TATGGGTTTC ANNAGGCACC NTTTTGCTAN 180
GNAGGCATAN TTANN 195

SEQ ID NO:6521

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07691

SEQUENCE DESCRIPTION:

GATCGAAACC AAGTTATTAT TATCCTTACT GTATTCCATT GCTTGATCA AAATCTACTT 60
TTTTNCTAGT AGGAAACTAT GCATTCCCTA TCAGTCCCTT TGATTATCAG GAATGTATAA 120
GAATTGTGAT AAAATAGGCC AATATGCTAT TTGGTACCAT TTGTATGTAT TTTANGTGCA 180
TAACCTAGGT AAAATAAGNN TAGTTTCTAT GTGTAAAGAA GCAAGTTTCA CAGTATGCAG 240
ACTCCAGTGG ATGTATTTTA GGCTGTATAC ATNTTTAAGC TCTNTNTGGT ANATAGGTA 300
TACAAACTAN NTGCCTCNA ATNCNAATAN NTTCCGN 337

SEQ ID NO:6522

SEQUENCE LENGTH:331

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07692

SEQUENCE DESCRIPTION:

GATCATTTCA TGGAACTTTT GAAGTATCTT TGAATCTAAC TTTGACTTGG TGGTGGACCT 60
TCCTTGTTTT TANTAACACC TAAGAGATAT CCTTTAGAAT TACATGTATT TTAGCATAAG 120
GAAATTGNAA AAGTAAAACA TACTGGTTTT TTTCAACAAG ACCATATGTA AATTAAATAG 180
TGAAATGTGT ATGNGTTTCA GTAGAACTGT ACCATCAACA ATGTTTCCAT AAATATGCAG 240
NGTTCTTNCT NNTGTATTGT TATTTACAAT ATTGTTAAAT NGGATGCATT TGCAATNNCT 300
CGGNTTCTAA AGNATTGNGG TNCNGANCGT N 331

SEQ ID NO:6523

SEQUENCE LENGTH:107

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07693

SEQUENCE DESCRIPTION:

GATCTGGAAT GGTGTTTTGG TTTGGGGGGA ATTTTTTTTT TTCCCGGCA AATNACATGT 60
GTTGTTGATG TGAGTATCTG ATGAAAAANC AATGNCAGAA TANCCGN 107

SEQ ID NO:6524

EP 0 679 716 A1

SEQUENCE LENGTH:196
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07694
 SEQUENCE DESCRIPTION:
 GATCTATATA TGTATAAATA TGTATTTTGT CAAATTTGTN ACTTAAATAT ATAGAGACCA 60
 GTTNTCTCTG GAAGTTTGT TAAATGACAG ANGCGTATAT GANTTCAAGA AAATTTAAGC 120
 TGCAAAANTG TATTTCTAT AAANTGAGNA GTCTCACTGN TAGCGGTTCT TTATTGCTCA 180
 TTTTTTAAAA ANTGGN 196

SEQ ID NO:6525
 SEQUENCE LENGTH:257
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07695
 SEQUENCE DESCRIPTION:
 GATCGGCTCT AGAAACACTG CTTTCCTTC TTTNCTTTTG TTTCTTTCT NTTATTATTT 60
 TTTTAAAGAG TAAACGAAAG TNCTGTATGA ATTCGGACCC AGGCAGCAAC CACAGGNCTN 120
 GGGGTGAGAC CCTCTCTCT CCAAGNGAGC AAAAAGGACC CATGGCCCC AAAAACC 180
 CTCAACGAGA ATCCTCAGCN TTGTAAATG CAAAANTGTC TAATGAATAT TTNTATNTGT 240
 NCCATTTTGT ATAANTN 257

SEQ ID NO:6526
 SEQUENCE LENGTH:297
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07696
 SEQUENCE DESCRIPTION:
 GATCCTATTT AACATGACAA GTAAATGCT TGCCATAGGA TATTTATATC TAGAACTTTG 60
 ACAGACATAC CTAATCATTG TCACTCAGGC ATCCTGCATA TCATTCATTT ATTCATTTGC 120
 CTATCATTTG TTGANCATTT AAAATGTCCT AGATACTTAG NATATGANTG TTGAATGTCA 180
 CAAAACTGAA TGTCACACTC CCTCCCCTTT AGANATTTNC AGCATGGGAT GCGAGCTAAT 240
 GATATGTAAA TAANTACATA TTTNTAANTA ACACAACCNN AAAANAANAN AAACNAN 297

SEQ ID NO:6527
 SEQUENCE LENGTH:137
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07697
 SEQUENCE DESCRIPTION:
 GATCGTGGCC AACGTGGACC TGCCTACGNA GGGTGGGCTC TGACCCAAGT GGGGCCTCCT 60
 TGTCAGGTC TCACTGCTTT NCACCGTGGT CAGNGGGACT NTCAGCTGAG CTTNAGCTCC 120
 CATGGAGCCA GCAGGNN 137

SEQ ID NO:6528

SEQUENCE LENGTH:293

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07698

SEQUENCE DESCRIPTION:

GATCAGCCTG AGAAGCCACT CAACTTGGCT CACATCGTGC CTCCCAGACC TGTAACCAGC 60
 ATGAACGACA CCCNTGTTCT CCCACTCTGT TCCCTCCTCA GGAAGTCCTT TTATTACCAG 120
 GAGCTCGGAC GGTGCTTTTG GTGGNTGCGT CTAGATGGAT AACATGACTT CTTCTACCCT 180
 AAAATATTCC TATAATACTT TNAGCTGTTC TGGTTCCTCC AGGGTGNATG GTACCNATTA 240
 ACCCAAAATA CTGNTTATNT ACCCNTTNTT ACNATNTANN AGTCANTTTT GGN 293

SEQ ID NO:6529

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07699

SEQUENCE DESCRIPTION:

GATCCACGGG GCCAGGCCAC CCTGCGGGAG CGCCACACGC ATCCACTTCG GATTCAGTGG 60
 GTGAAGACAG AACGCTNAGA GTCTGCAGGC GGCTCCTGTG CTTTAAATTT CTGGCTCTTC 120
 GGATGTCTTC TAGACATTTA CTATCACTGC ACCTGAAGAA AAAATCACTT TTACCTTCCT 180
 AATTTAAAAA GACAAAACAG AAATGTACGT TCCTTCGCTA GCTTTAGTCT TTCTGTTCCC 240
 ATTTTNATAA ANTCTGAGGC ATTGATAATG NTCNNATCTA AAATTTGTAC AGTGTGATTT 300
 CNNTNNTAAG NGTTAAANCT TTTANTAAAA GGGGCANGAN NTNAN 345

SEQ ID NO:6530

SEQUENCE LENGTH:266

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07700

SEQUENCE DESCRIPTION:

GATCCAGAAA ATAGGTACGG TTTTAAAATA TTTNATATAG AAAAGCTACA AAGTAAATTG 60
 AGCAATGCTT TTAAGTTAT CTTTGTTTTA TAGACTTTTT TGTGTATGT ATTACAGTCT 120
 TTATAATCTT ATTTAATGNA TATNTGTACT TTCAAGTACT GATGGAGATA GNCTCAAAAC 180
 AGTTATTTTN TTACAATTAA TCTACAANGG GCATTANTAT TGTNGNCTTT TAAACATCT 240
 GCTGGNTATC TNATNTGCAA TTCNTN 266

SEQ ID NO:6531

SEQUENCE LENGTH:43

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07701

SEQUENCE DESCRIPTION:

GATCCACAC ACTAAATATA TGGTTGTTTT GCCACCTTTT AAA 43

SEQ ID NO:6532

EP 0 679 716 A1

SEQUENCE LENGTH:285
SEQUENCE TYPE:nucleic acid
5 TOPOLOGY:linear
CLONE:HUMGS07702
SEQUENCE DESCRIPTION:
GATCTGTTTT TCACCACTTC ACTGAAAGAC ACCATTTATC TACCCAAGGG CAGAAAGTAG 60
AACTTACTAT TCATTAAATG TTTGACACAA TTGGAATTGT CTTTAATTTC TGTGAGAATG 120
10 CTATTGAAAA TGTGAATTGC ATGACTTGTA GCATATTCTT TTCTGCAAAA TAGCCATATT 180
ANCATGCTTA TGACAATGAC TGTGCTACTG TCTNTGGAAA ANTGTTTGTG TCAGTTGGNA 240
ATNCTAANNG NTTACCTGC GGANNAACG CCCCNNNCCN CCCAN 285

15 SEQ ID NO:6533
SEQUENCE LENGTH:43
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07703
20 SEQUENCE DESCRIPTION:
GATCCTCTTG TGTATATGCA AATAAACCTA TTATTCACTT AAA 43

25 SEQ ID NO:6534
SEQUENCE LENGTH:194
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07704
SEQUENCE DESCRIPTION:
30 GATCAAGACA CCTTTTATTA ATAAAGAAGN GACACAGGTG TAGATATGTA TATACAAAAA 60
GATGTACGGT CTGGCCAAAC CACCTTCCCA GCCTTTATGC AAAAAAAGGG GAGANTCAAA 120
GCTTTCATTT CAGAAAGTGT GCGTGGANAA GTATCTGTAA TTAAAGTTTC GANGTAATTT 180
AACCTATTTT TAAA 194

35 SEQ ID NO:6535
SEQUENCE LENGTH:45
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
40 CLONE:HUMGS07705
SEQUENCE DESCRIPTION:
GATCAAGCCC TGCTTAAAGT TAAATAAAAT AGAATGNATG ATAAA 45

45 SEQ ID NO:6536
SEQUENCE LENGTH:38
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07708
50 SEQUENCE DESCRIPTION:
GATCTTCTGG ATGACCTGAA ATNAGAACTA ATTGGAAA 38

55

5 SEQ ID NO:6537
 SEQUENCE LENGTH:52
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07709
 SEQUENCE DESCRIPTION:
 GATCGCCCAA CTGCACTCCA GCCTGGNAGC AAGACTCTGT CTCGAAAAGA AA 52

10 SEQ ID NO:6538
 SEQUENCE LENGTH:55
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS07710
 SEQUENCE DESCRIPTION:
 GATCAGNACC ACTGCACTCC AGCCTGGGCA ATCANAGTAC GACTCCTGGC TCAA 55

20 SEQ ID NO:6539
 SEQUENCE LENGTH:252
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07711
 25 SEQUENCE DESCRIPTION:
 GATCAGTCTG AAGTCGATTA CTAACAGCTG AAGCTGATAA AACCATTAAA GTATACAGAG 60
 AGGATGACAC AGCCACAGAA GAAACTCATC CAGTCAGCTG GAAACCAGAA ATTATCAAGA 120
 GAAAGAGATT TTAATGAATG TGGAAATTTT TCTCTCTCTT TTTTCTCTT TTTAATTAAA 180
 AAAAAAAGG CTTGGGGTTC ATGGGGTTTT CCNGTCATT NGTGCNCTGG CTGGGATTTT 240
 30 AAGGGGGAAT TN 252

35 SEQ ID NO:6540
 SEQUENCE LENGTH:48
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07712
 SEQUENCE DESCRIPTION:
 40 GATCGAGNCA CTACACTCCA GNCCGCCGAA GCAAGACTCT GTCTCAAA 48

45 SEQ ID NO:6541
 SEQUENCE LENGTH:235
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07713
 SEQUENCE DESCRIPTION:
 GATCANACAT TTNCTACCTA TGAATNTTGC TGCATACAGA AAGTGCCCTT TCCTCAGGAA 60
 50 GTTGCTGTGT NTCATTCTT TAGATGGNCT CTTATCTAGA ATACATAGCA GCTCTGCAAA 120

55

EP 0 679 716 A1

GAANCAGTTT TAAAAATGG GAACTTCTAC ATTGAAAAGT CCCNNTTTT GTGCCAACTA 180
TGATTAGTGN GCGGAAGNAN TCTTATTCTA TGGCATATGT ATGGANGGGG GGGGN 235

5 SEQ ID NO:6542
SEQUENCE LENGTH:36
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS07714
SEQUENCE DESCRIPTION:
GATCCTTTTA AATTAAAGC AAGTCCTAAA ATGAAA 36

15 SEQ ID NO:6543
SEQUENCE LENGTH:155
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07715
20 SEQUENCE DESCRIPTION:
GATCCAGCGN GCGCACGCAG AGTACCTGCA CAAGGCCGTC TTCAGGGGCC TNCTCACGGA 60
GAAGNCGGCG CCCGTNATGA ACGTCATCCA CAGCATCTTC AGCCTCGTGC TCAAGTTCCG 120
CAGCCAGCTN ATCTCCAGG CCTGGGGGCC CCNNN 155

25 SEQ ID NO:6544
SEQUENCE LENGTH:110
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07716
30 SEQUENCE DESCRIPTION:
GATCAAATAT CACTCTCCTA CTTACAGGAC TCAACATACT AGTCACAGCC CCTATAACTC 60
CCNCTTACAA TANNTANCCA CAACCAAAAT GGGGCTCANT CACCCAACCN 110

35 SEQ ID NO:6545
SEQUENCE LENGTH:192
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07717
40 SEQUENCE DESCRIPTION:
GATCTGGTTG TNTAGAAGCA TGCAGCACCT CCTGCTTCAC TCTCTCTGTC TCTCCTGCTC 60
CACCATGGCC AGAAACGTGC CTGCTTCCCC TTCGCCTTCT GCCGTGATTG TAAGTTTCTT 120
GAGGGCTCCC CAGCCATGCT TCCTGTACAG CCTGCAAAAC TGTGAGTCAA TTAAACCTCT 180
45 TTTNTTCATA AA 192

50 SEQ ID NO:6546
SEQUENCE LENGTH:73
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

55

EP 0 679 716 A1

CLONE:HUMGS07718
SEQUENCE DESCRIPTION:
5 GATCATACTG GTTTGTTTCT TATTTTGATT GGAGAAAAAT NAAATTGCTG CATTTCGCAG 60
TGACCCATTT AAA 73

SEQ ID NO:6547
SEQUENCE LENGTH:80
10 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07719
SEQUENCE DESCRIPTION:
15 GATCAAAATGT GTGGCCTTTG ACTGAAATCA GCCAGCCCAT GGCCCCCAAT AAAGGCAGCT 60
GCCTACTNCT CCNTCTGAAA 80

SEQ ID NO:6548
SEQUENCE LENGTH:115
20 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07720
SEQUENCE DESCRIPTION:
25 GATCCANATG CGTGTGGCAG CCNAAGGATT TGTGTAGGA GCAATGNCTG TTGGTATGGG 60
CTATTCCATG TATCGGGAAT TCTGGGNAAA ANCTAAGCCT TAGAAGAAGA GATGN 115

SEQ ID NO:6549
SEQUENCE LENGTH:98
30 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07721
SEQUENCE DESCRIPTION:
35 GATCATAGCA TATCTACTAT TTTATTATTT GATTTTAAAA ATTAAACATT ATATTATGGG 60
TAACCTTACA TGTCAATAAA CAATTCACACA TTGTCAAA 98

SEQ ID NO:6550
SEQUENCE LENGTH:33
40 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07722
SEQUENCE DESCRIPTION:
45 GATCAACTTT GGAATAAAGT GATGCCCTTC AAA 33

SEQ ID NO:6551
SEQUENCE LENGTH:42
50 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07723

55

EP 0 679 716 A1

SEQUENCE DESCRIPTION:
 GATCCACTGT AATATTTAAT AAAAAATGTT ACTATCTGTA AA 42

5
 SEQ ID NO:6552
 SEQUENCE LENGTH:24
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

10
 CLONE:HUMGS07724
 SEQUENCE DESCRIPTION:
 GATCAAATAT CACTCTCCTA CAAA 24

15
 SEQ ID NO:6553
 SEQUENCE LENGTH:141
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07725

20
 SEQUENCE DESCRIPTION:
 GATCAGGACC CTNTCAGTG CACCCANTTA GNGTTGTGAG CCCTTAAAG GACCAGGGAT 60
 TGCTCACTCG GGGAGCTCGG CTCTTGAGAC AGGAATCTTG CCCATTCCCC GAACGAATAA 120
 ACCCCTTCCT TAACTCAGAA A 141

25
 SEQ ID NO:6554
 SEQUENCE LENGTH:226
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07726

30
 SEQUENCE DESCRIPTION:
 GATCAGGCAG CAGGAGTGAG GGGCACAGTC ACCCCAGGCC TTGCTGAGCC CAGGTCTGGG 60
 TACTGAGTGT CCAGAGCTGC CTCCCAGGA GGTAAAGTG GGGGCAAAGG GGAAGCTTCA 120
 AGCACTTGC CTACTTTTGT TCACTNCCCA GTGCACTGTG ACTCAGGCCT TCCCATCAGG 180
 CCTATTTGTC TACCCAATAA AGCGTGTTTT TTCCAGAAAC AAGAAA 226

35
 SEQ ID NO:6555
 SEQUENCE LENGTH:46
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07727

40
 SEQUENCE DESCRIPTION:
 GATCAGCCTG GCCATGGTNA AACCCCATAT CTAGTAAAC TACAAA 46

45
 SEQ ID NO:6556
 SEQUENCE LENGTH:306
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07729

50
 55

SEQUENCE DESCRIPTION:

5 GATCCACTAT MAGGGTGGCA TCAGCCTGAA TTGAACATCC TATATCTNAA CATCTGCATA 60
 AATTCCAGGT GACCACCAGT GTGCACCACT CACCCATCCC AGTGTATTCA GGCCNNNTGA 120
 CCTGACCCAN TTTGGCCCAG AGGACTCTGC TGGTGACCTG AAAAGAGGGA CTACATATTT 180
 NAGGATATTG TTGGTCTCAC ATGACTCTTG ATAGAGTTTG CCAGCCTTCA AATCACCAAC 240
 TGTGGGAACC AAGNNGGNN TTATTGCCTT GCTGTGTTCA ATTACANAAA ACGNTGGAAA 300
 NGGCTN 306

10

SEQ ID NO:6557
 SEQUENCE LENGTH:255
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

15

CLONE:HUMGS07731

SEQUENCE DESCRIPTION:

20 GATCTGAATA CCCCTTCTCT GTAGCTGCTA GTGAGCCTTC CCATTTAGAT TAAAGATTGC 60
 TTTATCCAGC AGTCAATTAA CTCTCCAGTT ATCAGTACTC CCACAATTGG CCAGGGCAAC 120
 AATAATTGGA GTTCATACTG ATGCCCTGAG GCACTGAAAA AAAAAAAAAAT TCCAAGGNG 180
 CCTTNTGGGN TGNCTAAAGG TTNCATTGTG NTTGGNGGCT TTNGTGTTAG GTGTGCNGTT 240
 TANTTTCCTA NTTNN 255

25

SEQ ID NO:6558
 SEQUENCE LENGTH:59
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07734

30

SEQUENCE DESCRIPTION:

GATCACGNCA CTGCACTCTA GCCTGGGCGA CAGAGAGAGA GACTGTCTAA AAAAGGAAA 59

35

SEQ ID NO:6559
 SEQUENCE LENGTH:69
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07735

40

SEQUENCE DESCRIPTION:

GATCCTNAGG TCTCCTAGCC ATGTGGAGCT GTAAGTCCAA TNAAACTTCT TTTTCTTCCC 60
 AGTCTCAAA 69

45

SEQ ID NO:6560
 SEQUENCE LENGTH:53
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07736

50

SEQUENCE DESCRIPTION:

GATCTGTCAA GTCCAGTAGA GCTTCAAGGT AAAATNAAAA TAACAATGTG AAA 53

55

EP 0 679 716 A1

5 SEQ ID NO:6561
SEQUENCE LENGTH:33
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07738
SEQUENCE DESCRIPTION:
GATCCAGAGG CCATGGAAGA GTCTCCAAGG AAA 33

10 SEQ ID NO:6562
SEQUENCE LENGTH:158
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
15 CLONE:HUMGS07739
SEQUENCE DESCRIPTION:
GATCAGCTCA AAAAGNTTGG GCTTAGNGTT TTCATTGCAA ATNATAATTG CTGTAGAGCC 60
ACACACAAC TTTGAAC TTT TAATTATAAG TGTATGGCT AAAGTTATTN NCTGAAAATT 120
20 TCAGTAAAGT GTGTGAATGT TTCTTTATGT ATTANCNN 158

25 SEQ ID NO:6563
SEQUENCE LENGTH:121
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07740
SEQUENCE DESCRIPTION:
GATCAGGGTT CTTTTTTTCC CCCATACAAG TATCCAGTCA TTGTAACACT GTTTATTGAA 60
30 AGAATTATCC TTCTCTCATT AAATTACCTT GCCAATTAGT AAAAAATCAA TTAACCATAA 120
A 121

35 SEQ ID NO:6564
SEQUENCE LENGTH:87
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07741
SEQUENCE DESCRIPTION:
GATCAGCATA CACAAATNAC AAAAGTCTGA ATTTTTTTTT ATCAAGAGGG ATAAACACC 60
40 ATGAAAATAA ACTTGAATAA ACTGAAA 87

45 SEQ ID NO:6565
SEQUENCE LENGTH:54
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07742
SEQUENCE DESCRIPTION:
50 GATCCNTTTT NCTGTGGCAT TTGGTGCCCT AAAAAATTTG CATTTATCTT GAAA 54

55

EP 0 679 716 A1

5 SEQ ID NO:6566
SEQUENCE LENGTH:20
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07743
SEQUENCE DESCRIPTION:
GATCATTGA CAACTGCAAA 20

10 SEQ ID NO:6567
SEQUENCE LENGTH:143
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
15 CLONE:HUMGS07744
SEQUENCE DESCRIPTION:
GATCGACTAT GAGACAGCAA AGGGGCCTAC CTNCTNATGG AGTTAGCCAA GAGCCAGGAC 60
AAGAGTCTCA AGATTATGA CGGTGCCTAC CATGTTCTCC ACAAGNAGCT TCCTGACGTC 120
20 ACCANCTCCG TNTTCNTGA ANN 143

25 SEQ ID NO:6568
SEQUENCE LENGTH:199
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07745
SEQUENCE DESCRIPTION:
GATCTGCTGC AATGCTCNAA GGCCAAGGAT TCATCTNTCT TNTCACCGTA GGACCTGNCT 60
30 GGCATTGTAT TAGCAAATC TTCTACTAGA CATCGTACTA CACGACACGT ACTACGTNGT 120
AGCTCACTGC CNCTGATGTC CTATCAATAG GAGCTGTATT TNCCATCATA GGAGGCTTCA 180
TTCCTGGTN TCCCNATN 199

35 SEQ ID NO:6569
SEQUENCE LENGTH:37
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07746
SEQUENCE DESCRIPTION:
40 GATCGTTTCC AGGACAATAA AAGTTTCGGA GTTGAAG 37

45 SEQ ID NO:6570
SEQUENCE LENGTH:57
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07748
SEQUENCE DESCRIPTION:
50 GATCAGATGT TAAAAATNAT TCCAAATACT TCATTAAAAA AATTATTTTA TTATAAA 57

55

EP 0 679 716 A1

5 SEQ ID NO:6571
 SEQUENCE LENGTH:152
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07749
 SEQUENCE DESCRIPTION:
 GATCGAGGAA AGGGGCGGAG AATTTTTTTT TCTTCTTCCC ATTCTTTAA AAAACCAACA 60
 10 CAAGAAAACA CACACACACA CAACCAACGT TTGTTCCCGA GTAGAAACAT AAATAGGGCA 120
 GAATCGAACA CTCTGTTCTT CTCTCTTCCA AA 152

15 SEQ ID NO:6572
 SEQUENCE LENGTH:22
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07750
 SEQUENCE DESCRIPTION:
 20 GATCAGAGCT TGA AAAAACA AA 22

25 SEQ ID NO:6573
 SEQUENCE LENGTH:111
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07751
 SEQUENCE DESCRIPTION:
 GATCAACCTG TGGCTGTTTT CCCGTCTAGG TTCTCACAGG NATCTCCTGA CAGAGGTACT 60
 30 TAACAATGGC TCTNCTGGAA ATTTCTATAA ATAAAATGTC CAAAATGGAA A 111

35 SEQ ID NO:6574
 SEQUENCE LENGTH:44
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07752
 SEQUENCE DESCRIPTION:
 GATCTCATCA CTGCTCCTTA TAATAAACCT AATAAAGCAA GAAA 44

40 SEQ ID NO:6575
 SEQUENCE LENGTH:42
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS07753
 SEQUENCE DESCRIPTION:
 GATCCCGCCA CTGCACTCTA GCCTATGTGA CTCTGTCTCA AA 42

50 SEQ ID NO:6576
 SEQUENCE LENGTH:48

55

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07754
 SEQUENCE DESCRIPTION:
 5 GATCATTTGT AAGACCAAAA TATAAATAAA AAGTTTCAAA AATCTAAA 48

SEQ ID NO:6577
 SEQUENCE LENGTH:46
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07755
 SEQUENCE DESCRIPTION:
 15 GATCGATTAC AATAAATGTT TTTCTAAAAA GCCATTTTCC CACAAA 46

SEQ ID NO:6578
 SEQUENCE LENGTH:125
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07756
 SEQUENCE DESCRIPTION:
 GATCAAAAGC TTTGCATTGA TAAGTGTCTC ATAATATTG CTGTGATTGG AGAAAAAATG 60
 25 TAGTCGTAGC CAATAAATTT TATCAGCTTT TAAGTTTCAG TATTATTAAA CCATTTTCAT 120
 ATAAA 125

SEQ ID NO:6579
 SEQUENCE LENGTH:36
 30 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07757
 SEQUENCE DESCRIPTION:
 35 GATCAAAATGT GAAATAAATA TGAATGGAGT GGTAAG 36

SEQ ID NO:6580
 SEQUENCE LENGTH:199
 40 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07758
 SEQUENCE DESCRIPTION:
 GATCTCTGGC CCTTTGCGTT TCAGTATTTG TCCTTTTAT GCCTGAATGT TTAACCTGCC 60
 45 ACATNTTGT ACTGCAACAT GAATCTTGGA AATTTTAATG TTATGCATTT NAAANGTTTT 120
 TGGTTATCTG TAAACTAAAT GTGCCCTTAT TGGCTNGGNG GTCAAGTNAA GATGTTTGTA 180
 TAATGTATTN GTCAGNAAA 199

SEQ ID NO:6581
 50 SEQUENCE LENGTH:66

55

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07759
 SEQUENCE DESCRIPTION:
 GATCTACTGT GATGTTGTCT TCAAAGGCAG GNGAAAATAA TGTTACAAT AAAATGTGCT 60
 AACAAA 66

10 SEQ ID NO:6582
 SEQUENCE LENGTH:135
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07760
 SEQUENCE DESCRIPTION:
 GATCTGTAAA GAATGACTTT CAAGGATAGT TCTTAGGGGA CTGTGAAAGT GTTGGGTCTT 60
 ACTCNAGGAT GCCTGCATGG GACCCNACCC GGAGCTGGTG TGGCCATTCC CCAAGTGCCA 120
 CTGGNCCATG GNNNN 135

20 SEQ ID NO:6583
 SEQUENCE LENGTH:130
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07761
 SEQUENCE DESCRIPTION:
 GATCAGCCTG GCCAACATGA TGAGACCCCA TTTGTACTGG AAATGCAAAA ATTGCCAGG 60
 CATGGTCGCG GGTGNTGTG GTCCCAGCTG CTCGGGACGC TGAGGCAGGN NNNATCGTTT 120
 GAACCTNGGN 130

30 SEQ ID NO:6584
 SEQUENCE LENGTH:80
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07762
 SEQUENCE DESCRIPTION:
 GATCAAATGA AAGGCCTATA ACTAAAATCA GCCAGCCCAT GGCCNCACT AAAGGAACCT 60
 NCCTCTGCTC CCTNTGCAAA 80

40 SEQ ID NO:6585
 SEQUENCE LENGTH:48
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07763
 SEQUENCE DESCRIPTION:
 GATCAACAAT TTTGTTTTG ATTAATAA GTTTTTGAA AGTTGAAA 48

50 SEQ ID NO:6586

55

EP 0 679 716 A1

SEQUENCE LENGTH:178
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07764
 SEQUENCE DESCRIPTION:
 GATCCCCTGC TGGCTGGGGG CAGCTCCCAG GATATACTGC CTTACAACCTG TTTCTAAAGC 60
 CCATCCTNCT AACATGGCGA TTCCGGAGGT CAAGGCCTTG GGCTCTNCCC AGGGTCTAAC 120
 GGNTAAGGGG GCCACATACC NGTTCCAAGG GGNNTGTCAA GTNGTGAAGT CGTTNTGN 178

SEQ ID NO:6587
 SEQUENCE LENGTH:99
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07765
 SEQUENCE DESCRIPTION:
 GATCAGATGG TTTATAAAAG TAATAACCAT AAAGCAAAAA ATAATTGAA AGCCCGTCTA 60
 TTCCTATGCT CANTAAAGTT AAGGTTTTTC TTCATTAAA 99

SEQ ID NO:6588
 SEQUENCE LENGTH:65
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07766
 SEQUENCE DESCRIPTION:
 GATCCTTTGA GCCCAGGAGT TTGAGACCAG CTTGGGCAAC ATAGTGAGAC CCTGTCTCTA 60
 CCAAA 65

SEQ ID NO:6589
 SEQUENCE LENGTH:205
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07768
 SEQUENCE DESCRIPTION:
 GATCACAGGG TCAGGAGTTC AAGACCACCC TGGACAATAT TGTGAATTCC TGCCTCTACT 60
 AAAAATACAA AAATTAGTCG GGCATGGTGG CGGGTGCCTG TGGTCCCAGC TACTCGGGAG 120
 GCTGAGGCCA GAAATCGGT TGAACCTGGG AGGTGGAAGG TGCACTGAGC CAATATCACA 180
 CCAGTGCACCT CCAACCTGGT GACGN 205

SEQ ID NO:6590
 SEQUENCE LENGTH:57
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07769
 SEQUENCE DESCRIPTION:
 GATCTCATAC TCAACAGAAG GAGACTCAGA GACTCCAGAA GGCACAGGAG CTGCCCN 57

SEQ ID NO:6591

SEQUENCE LENGTH:184

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07770

SEQUENCE DESCRIPTION:

GATCATGTTT ATTTACTGAT AAAACTCAAT ACCTAGAACT GCACCTAGCT CATGATGGTA 60
 CCTAATAAAT ACTTCAATGA AATGATGTGG TGATGCATGC CTATAGTTT AGCTACTTGG 120
 GAGGGTGGCC TGAGCCCAGG GGTCTAGAC CAGCTTAGGC AACATAGCAA GACCCCATCT 180
 CAAA 184

SEQ ID NO:6592

SEQUENCE LENGTH:165

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07771

SEQUENCE DESCRIPTION:

GATCTGAATG GTTTAATAA GTGCCTGGNA TTCCCTAC TGGCTCTCAT NCTNACTCTT 60
 GCCGCCCTGT GAAGAGGTGC CTCCACCGT GATTGTNAAG TTCCTGAGG CCTCCCAGC 120
 CATGTGGAAC TGTGAGTGA AAATNAAACC TCTTTATAA TAAA 165

SEQ ID NO:6593

SEQUENCE LENGTH:134

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07773

SEQUENCE DESCRIPTION:

GATCCAAGCT TAATTTTCAA ACAGCTCCC AAAAGCTGT ATCCTGAATA TCATAAAGTC 60
 TATCATATGA TGTGTACTCA GTCATATTCT GTAGACTCAA ATCTTTTATC TGATTATACC 120
 AGCTTCCCG NAAA 134

SEQ ID NO:6594

SEQUENCE LENGTH:148

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07774

SEQUENCE DESCRIPTION:

GATCTGCCCC CGTTGGCCTC CAAAGTGCTA GGATTATAAG TGTGAGCCAC CACACCAGGC 60
 CCGTTTCATC CCTGNCTCAG CCGCATGCC AAGCTCCCTN GGTGCCCCGNC CACAGGTGTC 120
 ACCCTTTNAG GAGAANGTGG CCTGGGGN 148

SEQ ID NO:6595

SEQUENCE LENGTH:238

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS07775

SEQUENCE DESCRIPTION:

5 GATCGCTTGT AGCTGGGAAC AGTGCTTACT TCAGGGAAAA CTGATTTTGA TAGAACCGAC 60
TAGTTGAGTA TTAAACTGTT TTGGAAATAA TTTATATATC TAAGTGCTGA AATATATTTC 120
TCTTCTCCCC ACTTATTCAA ATAAAAANGT TTTNATACTA TAAACTTTTC NNNGTTAGCG 180
CCACGCAGTC CTTGTCAAGN TTGTNCTGGG GGNCAANAAC TTGATGTATA AAGCATGN 238

10

SEQ ID NO:6596

SEQUENCE LENGTH:102

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

15

CLONE:HUMGS07776

SEQUENCE DESCRIPTION:

GATCTCTACT ACCANAAGGA AAATAGTTTA GGNGAAACCA GCTTTTANTG TTTTGTAAA 60
ATTAGCTTCA CCCTGTNATG TTANCAAGGT ATGTCTTTGC CN 102

20

SEQ ID NO:6597

SEQUENCE LENGTH:65

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

25

CLONE:HUMGS07777

SEQUENCE DESCRIPTION:

GATCGCGCCA CTGCACTCTA GCCTGGGTGA CCAGCCTGGG TGACAGAGTG AGNCTCCGTC 60
TCAAA 65

30

SEQ ID NO:6598

SEQUENCE LENGTH:282

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35

CLONE:HUMGS07778

SEQUENCE DESCRIPTION:

GATCTAGAAG AAAGTATTGT ACGTTTGAAT GCAGATTTTN ATCCACAGAT AGTTGTAGTG 60
TTTAGACATG ACAGGACCTA TCGTTGAGGT TTCTAAGACT TACTATGGGC TGTAACCTG 120
TTTTTNAAAA CTATTTTGA AACCTGAGAC TTGCCGTCTG GCATTTNAGT TTAATACAAA 180
40 CTAATGATTG CATTTGNAAG NGNTNCTTGA CCTATTCTTCT AAACGTCTAG NGCTCTGAAA 240
TGTCTTGCTG GNNGGTATTA AACTATTTGC CTGTNGTACA TN 282

SEQ ID NO:6599

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07779

SEQUENCE DESCRIPTION:

50 GATCAAATGG CTTAGAGNTA GTCCTGGAAT ATTCATATTC AAAGATTCCT TATTAATGAA 60

55

TGTCITTAAC TTAAATCTAC CCAATAATTG CAACATGGTT CTTTGTACAT TTTCATTATA 120
 TGGTGTTAAC AAGCTTCACT GCAAACAAAT AAATTACTTA AGTTATTTGN AAA 173

5

SEQ ID NO:6600
 SEQUENCE LENGTH:263
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

10

CLONE:HUMGS07780
 SEQUENCE DESCRIPTION:
 GATCAGTGTA GCAAATNAGG GATGAAGNGT TCAAACCTTT TGGCCCTTTC TTTCTTTNCT 60
 AGGCTTCTCC CTCGAGGGT GTTCCGTAGT TTCTTCTCGA GCCAATGCAT GTATTATAGC 120
 AGCAGGTGTC TTTGTGCTTT CTCATCATAG TAACGTACTA CTTGTAAATA CATTNTNCTA 180
 TTTNCTATTT TTTTGTATTT TTTTGTGACA TTTTGTTCCT TTGGTGTGCT GNGNGNNNCC 240
 CATGCCCTCA CTCCTTTAAG AAA 263

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SEQ ID NO:6601
 SEQUENCE LENGTH:51
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07781

25

SEQUENCE DESCRIPTION:
 GATCCAAAAT TCTCAACAAA ATATTAGCAA ATTGAACCCA CCAAGGTAA A 51

30

SEQ ID NO:6602
 SEQUENCE LENGTH:283
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07782

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SEQUENCE DESCRIPTION:
 GATCAAAGGC CTCTNAGGGG TGTCCACCAA AAACCTCTCC TTCAAAGAG AAGACTCCGT 60
 GCTTCAGGGC TATGACATCA ACAGCCAAGG GGAAGAGTCC ATGGGAAATG CAGAGCCCCCT 120
 TAGGAAGCCN ATCAAAAACC GGNGCNTANN GTTAANGAAA GTGAAGTCCC AGGAAATACA 180
 CATGCTCCCA ATCAAAAAC AACGGCTGGC CACCTTTTTT CCAAGAAAGT AAATAACGGC 240
 TTTTAAANT NNGTATGNTT ATAATATGGG GAAAGGTGCA TTN 283

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SEQ ID NO:6603
 SEQUENCE LENGTH:223
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07783

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SEQUENCE DESCRIPTION:
 GATCGTTCAT CCGATTGCNA CGTTCCAAAG TGTCCAGGTT TTTAAGACCT TACAAATGAC 60
 TCAGCCCACG TGCCACTCAA TACAAATNTT CTGCTATAGG GTTGGTGGGG CAGAGCTGTC 120
 TTCCTTCTGC ATGTCAGCAC AGTCGGGTAT TGCTGCCTCC CGTATCAGTG ACTCATTAGA 180
 GTTCAATNNN NATAGATAAT ACAGATATTT TGGTAAATTG AAA 223

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5 SEQ ID NO:6604
 SEQUENCE LENGTH:100
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07785
 SEQUENCE DESCRIPTION:
 GATCACCTGA GCCTGAGAGT TCCAGACCAN CCTGGGCAAC ATAGCGAAAC TNCATCTCTT 60
 10 TAAAAATAAA ATAAATTA AATTAATACA AATAATTAAA 100

15 SEQ ID NO:6605
 SEQUENCE LENGTH:69
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07786
 SEQUENCE DESCRIPTION:
 GATCCCTTCC ACTCAGGATT CTCGCCGCT TTTCTAAGAA AATAATAAAA AAAAAATNCT 60
 20 TGTTTCAAA 69

25 SEQ ID NO:6606
 SEQUENCE LENGTH:164
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07787
 SEQUENCE DESCRIPTION:
 GATCTGCCCTG CCTCGGCCTC CCAAAGCGCT GGGATTACAG GCGTGAACCA CTGCGCCCAA 60
 30 CCTTTGGATT ATTTTGTGA GAGATGGGGC TTCCCTATGT TGGCNAGGCT GGTCTCAAAC 120
 TCCTGGANTC CAGCCTGGGC GGCAGAGCAA AACCTTGTCT CAAA 164

35 SEQ ID NO:6607
 SEQUENCE LENGTH:73
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07788
 SEQUENCE DESCRIPTION:
 GATCACTCCT CTGCTTCAGC CTGGGCAATT GAGCCAGACT CTGTCTCAA AATAAACAAA 60
 40 AAAACTTGAC AAA 73

45 SEQ ID NO:6608
 SEQUENCE LENGTH:74
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07789
 SEQUENCE DESCRIPTION:
 GATCTTTTGA CCTGGATATG TCTGTAAGGC TCCTGAAAGA GACAAATAAA GTCAATATAT 60
 50

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EP 0 679 716 A1

TTNCACAGTG CAAA 74

5 SEQ ID NO:6609
SEQUENCE LENGTH:58
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07790
10 SEQUENCE DESCRIPTION:
GATCTTAGGT CAAATAACAG ACTTGAGAAT ACTTTATAAA CTGCCATGAT ACAGCAAA 58

15 SEQ ID NO:6610
SEQUENCE LENGTH:139
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07792
SEQUENCE DESCRIPTION:
20 GATCGNGNGC GTTTAACGCT GGTGGGCAGC AATAAGGGGC AGATGTGACC AGATGCCTGC 60
NTCCCCAGGG TGCCGAGGGC AGCAGGAAAA AGTGGGGACC TCGGTGCATT TGCCCCACCC 120
NTCCCCTNNC TGGGCTAAN 139

25 SEQ ID NO:6611
SEQUENCE LENGTH:75
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07793
30 SEQUENCE DESCRIPTION:
GATCATCGAC ACTTCGAACG CACTTGCGGC CCCGGGTTC TCCCGGGGCT ACGCCTGTNT 60
NAGCGTCGCT TGAAG 75

35 SEQ ID NO:6612
SEQUENCE LENGTH:171
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07794
40 SEQUENCE DESCRIPTION:
GATCTGGGGG AGGAAGTTCA AGGAAGGCAC CACACCAGAA ATNGGACCTA AATTAAGGCT 60
TAAAGAATGA GCATGGCCAC ACTGTCAGTN AGTGTCTTT AGGTGGTAGG ACTATGGTTG 120
ANATTTTCNT TCTTCCTATC TTCCTGCCTT TTTCAGGNNN TNAATATTAA N 171

45 SEQ ID NO:6613
SEQUENCE LENGTH:69
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07795
50 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

GATCACGATG TAAATAATTA AGCATTTTGG TTGGATTCT TTTGTAATAA ACTATTTTGG 60
TACCATAAA 69

5 SEQ ID NO:6614
SEQUENCE LENGTH:271
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07796

10 SEQUENCE DESCRIPTION:
GATCTAGGAT AGCTAAGTNA CCTACATGGT GACTGTTCGG TCTAATTCAG AACAGAAAGT 60
ACTAAAAGAA TACTAAAAAG TTTTACCCA TTATCTCTAT CCCTGTTTNN TTTTGCAAT 120
TCTCAGCNGN AGGGAGCAAG AAAATACACA GCAGNGGGGG GAATGTTTGT TTGCATCCTG 180
15 GCTTTCTCAA CCGCACTTTA CGTCCTTGAA TATNCTAGAT ATTTACTCCT TCAAGAAGTC 240
AGGAGTATAT CCAGGCCTAC CTATCANCCT N 271

SEQ ID NO:6615
SEQUENCE LENGTH:237
20 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07797

25 SEQUENCE DESCRIPTION:
GATCACACCA TGGAGACTTT CTACTAGAGG ACTTGAAAGA GAACTGAGGG GCCACAAAAT 60
AAACTTCACC TTCCATTAAG TGTCAAGCA TGTCTGCAAA TTAGGAGGGN GTTAGAAACA 120
GTCTTTTCA TCCTTTTGA TGAAGACTGA AATTGTGCCG TGTGCTTA TATGANTATG 180
CAGTATGGGA CTTTGAAATA ATGATTCATA ATAAAATACT AAACGTGTGT CTTCAA 237

30 SEQ ID NO:6616
SEQUENCE LENGTH:196
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07798

35 SEQUENCE DESCRIPTION:
GATCTNAAAT AAAATTCAAA CTCTACTTCT GTTGTCTAGT ATGTAATTGA GCTAATGATT 60
CATTGGGAAT ATACCTTTTT ATACTCCTTT ATCATGGTAC TGTAAGTGA TCCATATTAT 120
AAATATAATT ATCTNAAGGN TTTTNAATTT TTNTNATGT CCAAGNGCCC ACGTGAATTT 180
40 NCTGGTGAAA GTNAGN 196

SEQ ID NO:6617
SEQUENCE LENGTH:75
45 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07800

SEQUENCE DESCRIPTION:
GATCATCGAC ACTTAGAACG TACTCCNGNC CCCGTGCTCC TCCAGAACCT ACGNCTGCAC 60
50 TAAGCGTCGA TTAAT 75

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EP 0 679 716 A1

5 SEQ ID NO:6618
 SEQUENCE LENGTH:50
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07801
 SEQUENCE DESCRIPTION:
 GATCACGCCA CTGTACTCCT GGGCAACAAA GCGAGACTCC CTGTCTCAA 50

10 SEQ ID NO:6619
 SEQUENCE LENGTH:208
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS07802
 SEQUENCE DESCRIPTION:
 GATCGAGGCT TGTNATGCCT TGTGAGAAAT AAGGGCCTTA ATTTGTACTG TCTGCGGCAT 60
 TTTTATAAT ATTGTATATA GTGACTGACA AATATTGTAT TACTGTACAT AGAGAGACAG 120
 GTGGGCATTT TTGGGCTACC TGGTTCGTTT TTATAAGATT TTGCTGGGTT GGTTTTTTTT 180
 20 TTAATTAATA AGTTTTGCAT CTTTTAAA 208

25 SEQ ID NO:6620
 SEQUENCE LENGTH:151
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07804
 SEQUENCE DESCRIPTION:
 GATCAGGAAG CCAAACCTAA GAAAGAGCTT GAAAATACTG CTCTAGAGCT TAGTCAGAAA 60
 30 GAAAAACAGT TTAATGCCAA AATGCTGGAA ATGGCACAGG CTAATCAGC TGGAATCAGT 120
 GATGCAGTGT CAAGACTGGA AACAAACCAA A 151

35 SEQ ID NO:6621
 SEQUENCE LENGTH:33
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07805
 SEQUENCE DESCRIPTION:
 40 GATCTNGGCA ACACAGTCAG ACTCCGTCTG AAA 33

45 SEQ ID NO:6622
 SEQUENCE LENGTH:321
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07807
 SEQUENCE DESCRIPTION:
 50 GATCCGGACG ACGTAATCAC TGACATTNAG ATGGTGGACG AGGGCTGGTG GCGGGGACGT 60

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5 TGCCATGGCC ACTTTGGA CTTCCTGCA AATAATGTCA AGCTTCTGGA GTNACTAGAG 120
 CTCACGTGCT ACTGCAACTN TAATTTCCCA TGTCCAAAGN GGTCTGCCT CCACCCCTC 180
 CCTATTCTG CTGCAAAATNT CTAACCAGAT GAGGTTCTGG ACAGACTTCC CTCTCCTGCT 240
 TCATTAAGGG CTTGGGGCAG AGNCAGCATG GGAAGGAGG TCCCTTCCC CAAGAGTCCT 300
 CTTCTATCCC TGGATGAANN N 321

10 SEQ ID NO:6623
 SEQUENCE LENGTH:96
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07808
 SEQUENCE DESCRIPTION:
 15 GATCCCAGCA CCCATTCTT AGCTATTTAT CCCTTCTG GTTCCGAAA GGCAATTATA 60
 TCTATTATGT ATAAGTAAAT ATATTATATA TGGAAA 96

20 SEQ ID NO:6624
 SEQUENCE LENGTH:242
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07810
 SEQUENCE DESCRIPTION:
 25 GATCTAGGCT TCTNGCCTCA CTCCCATTAG GAATAACACC CAGCAAAATG GAACTATCCA 60
 AGCACTTCTT ATACATCTGT CTCCCATTG TCTCTTAAGA AGTTTCTTCA GCTGGTGCGC 120
 CATCATCATC ACCATGACTG CCCCAGCTC TTGAGGAAAG GNATAATTAC CTCCATGGTG 180
 TTGCCTCCTA TTTGCCTAAA GCACCAGAAA TAATATTAATA TACACCAGAT TAACTACANA 240
 AA 242

30 SEQ ID NO:6625
 SEQUENCE LENGTH:97
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS07811
 SEQUENCE DESCRIPTION:
 GATCCCGTCC TTGTCAGAGA AGTGTTCCTT GGATGAGCGC CCAGAGGGAG GAGCCCATGG 60
 CCCTTCTTC TGTNACTGAT TAAAAATCTC GATGAAA 97

40 SEQ ID NO:6626
 SEQUENCE LENGTH:98
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS07812
 SEQUENCE DESCRIPTION:
 GATCCCTTCA AGACTTTGTC ACAGCCTCTA TCACACATCT GTTTTCTCG AAGNAAAAA 60
 TATAATTAAT AAAAATGTTT TACTCTTTTA CACTGAAA 98

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5 SEQ ID NO:6627
 SEQUENCE LENGTH:36
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07813
 SEQUENCE DESCRIPTION:
 10 GATCCCGNTG TTCTTAATAA ATCCTGACTT CCCAAA 36

15 SEQ ID NO:6628
 SEQUENCE LENGTH:35
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07814
 SEQUENCE DESCRIPTION:
 GATCAAAAAT AAATACTGCA CATAAANAA AAAAN 35

20 SEQ ID NO:6629
 SEQUENCE LENGTH:77
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS07815
 SEQUENCE DESCRIPTION:
 GATCTTGTTT TTATTGTTAT TGCTTTTATT ATNATTGCTT TTATTATCAT TAAAACTCTA 60
 GTTCTTGTTT TGTCAA 77

30 SEQ ID NO:6630
 SEQUENCE LENGTH:98
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS07816
 SEQUENCE DESCRIPTION:
 GATCCAGTAC AGAAAGGAAA GGNTATTTAT TGATTAACAG AAGTTGTCTT TTTAAAAGTG 60
 TTTATTTTGG GCAATAAAGA GCACAACATA ATCTCAA 98

40 SEQ ID NO:6631
 SEQUENCE LENGTH:132
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS07817
 SEQUENCE DESCRIPTION:
 GATCAGGGAA CAAAATCCTC TCCTTGTTGA AATATCCCAT GCAGTTTGTT GATACAACTT 60
 AGTATCTTAT TGCCTAAAAA AAAATTNNCT TATCNTTGTT TCAAAAAAGC AANNTCATGG 120
 ANAATTTTGG TN 132

50 SEQ ID NO:6632

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SEQUENCE LENGTH:304

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07818

SEQUENCE DESCRIPTION:

GATCAACGGC TGACCTTCTG GCGTCTGGGG CATGGTGAAC CCACCTNCAT GAATAGCACT 60
 GTGTTCCATG TGCCTGANGT GGCTGACATG GACTGCTGGC CTGTAAGCCC TGAGTTTGGC 120
 CACCGTTGTG CCCTTGGGGG TCAGGGGNTT NAGGTTTACA ACTGGTATGA CTGAGGTATC 180
 CTGCGGTGGC TGGCGTGCTG GGCATGGGGC NTGCTCACAG ACAGNATGGA GCAGGGAAGG 240
 GCTGTCCTGT GCCCATGCTC AGCTTNCCT TNNNGNNGAN GGAGGTNGTG GCCGTGGGTN 300
 CCTN 304

SEQ ID NO:6633

SEQUENCE LENGTH:35

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07819

SEQUENCE DESCRIPTION:

GATCTCAAAT AAAATTNCTG TAAAGAATTT CCAAA 35

SEQ ID NO:6634

SEQUENCE LENGTH:185

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07820

SEQUENCE DESCRIPTION:

GATCCAAATC GGGCCATCTT CATTGCCATG AGTGTTCGGA CAGGCTCTGC GTGCCAGGCA 60
 ACAGTGGGCA GACAAAGGGC TCCCTGGCTA CTGGGAAGAT ACAGAACCAT CCTGTGAAAA 120
 TCATTTCAGC ACTGGAGACC TTCTTTGCTA TTACTTCTTT CATTAACAAT GACCTCTAAT 180
 TTAAA 185

SEQ ID NO:6635

SEQUENCE LENGTH:151

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07821

SEQUENCE DESCRIPTION:

GATCCGTGGC CTGAAGGTTT CTAGAGNCGT CAAGAAATNA ATATCTTACA CTGTGATTCT 60
 GTGAGGAAAG ACTGGTAACC CAAACTCTC TTCTCTAATG TATTTTNAAC CGNAAATGAC 120
 AATATTTCTT TAATAAGTA TTTATACCAA A 151

SEQ ID NO:6636

SEQUENCE LENGTH:165

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS07822

SEQUENCE DESCRIPTION:

5 GATCCAGGTC TAGGCCCTTN ATCTGCTGCT CTGTGGCCCA GGGCAGGTTT GCTTGACCTC 60
TGCCTCAGTT CTCGACTCTA AAGGACATAC TGACCTACCT CACAGGGGTG TTGTNAGGAT 120
TAATAAATNT TGGTACTCTG CTTTGGAAAT NTGAAAAAAA AGAAA 165

SEQ ID NO:6637

10 SEQUENCE LENGTH:71

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07823

SEQUENCE DESCRIPTION:

15 GATCAAATTC AGTGTGTAT GTATGTTTAT GGCAACATTT TGAAATATAA AATTAAATGT 60
TTTNCCTCAA A 71

SEQ ID NO:6638

20 SEQUENCE LENGTH:348

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07824

SEQUENCE DESCRIPTION:

25 GATCTCAATA CTTCCCTGAC ATCCCTTAAG ACTCAAGTCT TGATACTTTG TCTGCTCTCT 60
GGCTCTTTCT TTGTCATTAC TATNATNNA ATCTTCTTGA TGTGACTCT TGCTAAAATT 120
TTTNGTCTTC CTTATTGTTA ATGCTTATAG AAAATGATTC CTTGTTATCT TTNAGCCCAT 180
GTATCCGAA TCATTGCTG TTTCTAGCCT AGGTGCTTTG ACTATGCCAG CTTGTATTTC 240
TNCAGAAATTT NCATTANCTA TATGTGTGGT ATTGTAGGCA TTACCATATT TNAATAATAA 300
30 NTTTAGGCCA TTGCTTCAA TCACAACAAA NNACAACNCA ACNNACNN 348

SEQ ID NO:6639

35 SEQUENCE LENGTH:53

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07826

SEQUENCE DESCRIPTION:

40 GATCTGAAGA GTGAAAATAA TGCTTTAAAT NAAAAATTGA AATCAGAAGG AAA 53

SEQ ID NO:6640

SEQUENCE LENGTH:127

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS07827

SEQUENCE DESCRIPTION:

GATCTTTTIN AATTCAGTAC AACCCATAAT CATGTAAATN CTCATTTNCT TTAGGNCATA 60
AAGAGAGCCC TAGGGTGCTC TGAATGTGTA CATGTNCTTG TCATAAAATG CATACTGTTG 120
50 ATACAAA 127

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5 SEQ ID NO:6641
 SEQUENCE LENGTH:23
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07828
 SEQUENCE DESCRIPTION:
 GATCTGTTGT TACTTAACAG AAA 23

10 SEQ ID NO:6642
 SEQUENCE LENGTH:85
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS07829
 SEQUENCE DESCRIPTION:
 GATCTTNATG TTAATNTTCA GAGGACTCTA GACAACAACT TAGCAACTGA AGCTTATNAA 60
 AGAAGAATTA AGCGCCTTGA GCAAA 85

20 SEQ ID NO:6643
 SEQUENCE LENGTH:113
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS07831
 SEQUENCE DESCRIPTION:
 GATCTTTTGT GGATATTTGT CACTTGGTTT CAGAAAGTNC ACAAATGTAG CAACAGCTCA 60
 CATGACTGAG TAGGTAGAAA ATGTGAAATA AATCTCATAT ATATNGTTTT AAA 113

30 SEQ ID NO:6644
 SEQUENCE LENGTH:54
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS07834
 SEQUENCE DESCRIPTION:
 GATCGTCCCG CTGCACTCCA GCCTGGCCAA CANAGCGAGA CTGTCCCCC CAAA 54

40 SEQ ID NO:6645
 SEQUENCE LENGTH:59
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07835
 45 SEQUENCE DESCRIPTION:
 GATCTTGCTC CTCTGTAAGG AACAGGGTG CCTAATAAAT ACATTNCTCC TTIATTAAA 59

50 SEQ ID NO:6646
 SEQUENCE LENGTH:184

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EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07836

SEQUENCE DESCRIPTION:

GATCCAGGGT CCCCAGGGT GGGCTCAGCT CCAGGGAGAG GCCACCCACG TGGCAGCCCC 60
ACCTCTTGAG AGCCCCAGT GCCGGAGCAG AAAGGACCCT GGACCCNGAG GCAGNNACTG 120
CGGGGTGGTA GAAAAGGTAG AGTAGGCTGT GGCAATGGAA TAAANCACGA TAAAAACGT 180
TAAA 184

SEQ ID NO:6647

SEQUENCE LENGTH:166

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07837

SEQUENCE DESCRIPTION:

GATCCTTCCC CTGCAACCAG ACAGTCTACA ACTGCCCCCT CCAGCCCATT TNCTGCCGTG 60
AAACCCACG CAGCCANACC AGGCTCTGGA ACCCTTTTTC GACTAGGNCA ACTCTTGGNC 120
ACCAGGNGGG GTAGAACC NAACACCAA CTGTACAGAN TNTCCN 166

SEQ ID NO:6648

SEQUENCE LENGTH:227

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07838

SEQUENCE DESCRIPTION:

GATCTNTNAT GGTACGACC ACCCGCNTAC CCAGTTCCTC CAGTGGAGAC CAGTTGGGTA 60
GTGGTGACTG GGTACGCTAC AAGCTCTGCA TTGTNTGCTG ATGGGACGCT CTTCAAGGTG 120
CAAGTAACCA GCCAGAACAT GGGCTGCCAA ATCTNAGNCA ACCCCGGTGG AAACACTNAT 180
TAAAGAACC CCAGAGGACA CCAACAGGTG TNTATCTGGN CNTGNCN 227

SEQ ID NO:6649

SEQUENCE LENGTH:314

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07839

SEQUENCE DESCRIPTION:

GATCTTTGGG GTTGGAGTGA AGGTCTTATC TGCATTAATT TTGTTAATGC ACATTGTAGA 60
TGTGGAGAAC ATCTTGACCT GCATAAGGAT TGCCTCTCAT TTTATTGTGT AGTGCTTTTT 120
CGAAATATTT ACATTTCTTT GGTTCCTATG TTCATGTATT CTGGAGAAGT AAAATTCTTT 180
GATTACTGGG CAGACCCTAA AACTACTAAG AGAAACATAA CATGTAGTTA GCTGTCTGCT 240
CATGGTTTGT TCAAATCTAA GCTAACCTTA ATGGTGCTGT GTCCGCTGTN CACAGAAAAC 300
GGCCTTAAAG TAGN 314

SEQ ID NO:6650

SEQUENCE LENGTH:104

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07840
 SEQUENCE DESCRIPTION:
 5 GATCCCCCTCC CCTGCCCCTG GGTTCCTGCC TTCNTCCTC AAGCAGGCAC NNAGGCTTTA 60
 GAGAAGTATA GGGGGCTTCT CCCCTGCNGG GCTTACCACA NTTN 104

SEQ ID NO:6651
 10 SEQUENCE LENGTH:358
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07841
 SEQUENCE DESCRIPTION:
 15 GATCAAGATG CCGAACCCCTG TGTCAGAAC CTCCAAATAC TGCCATGAGA AACTAGAGGG 60
 CAGGTCTTCA TAAAAGCCCT TTGAACCCCC TTCCTGCCCT GTGTTAGGAG ATAGGGATAT 120
 TGGCCCCCTCA CTGCAGCTGC CAGCACTTGG TCAGTCACTC TCAGCCATAG CACTTTGTTC 180
 ACTGTCTCTGT GTCAGAGCAC TGAGCTCCAC CCTTTTCTGA GAGTTATTAC AGCCAGAAAAG 240
 20 TGTGGGCTGA AGATGGTTGG TTTCATGTTT TTGTATTATG TATCTTTTGG TATGGTAAAG 300
 ACTATATTTT GTACTTAACC AGATATATTT TTACCCGAGA TGGGGATATT CTTTGTAN 358

SEQ ID NO:6652
 25 SEQUENCE LENGTH:46
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07844
 SEQUENCE DESCRIPTION:
 30 GATCCAGTTT GGGATTATGA AATAAACCAC AAATNAAAAT TTAAAA 46

SEQ ID NO:6653
 SEQUENCE LENGTH:63
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07845
 SEQUENCE DESCRIPTION:
 GATCTGGGGG GATGGAAAGC AAAATAAAGG AGGTGCCTGG NTGCATTATT TGCAGTGGGC 60
 40 AAA 63

SEQ ID NO:6654
 SEQUENCE LENGTH:290
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07848
 SEQUENCE DESCRIPTION:
 GATCCACCCA CCTCGGTCTC CCAAAGTGCT GGGATTACAG TCTTGAGCCA CCGGGTGACC 60
 50 GTCTTACAGG GATATTTTNA ATCCCGTNAT GGACTCTGTC TCCAGGAGAG GGGTCTATCC 120

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ACCCCTGCTC ATTGGTGGAT GTNAAACCAA TATNCCTTTC AACNGGTGNC TGNTAGGGAA 180
 AANCTACTCC TCATTATCAT CATTATTATT GCTCTCCACT GTATCCCCTC TACCTGGCAT 240
 GTGCTTGTA AGTNCTAGTT GTTCAATAAA TTTGTNAATA ATGCTGTAAA 290

SEQ ID NO:6655

SEQUENCE LENGTH:145

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07849

SEQUENCE DESCRIPTION:

GATCTTCTG GAAATNAGCA GTGACTAACG CTCACATAAC TGGTTTTTTT TTTNAACTGG 60
 GCTGATGAAT ACATTTACCT ANGAACTCA TTTCGTTTTA CTTAAGAGGG GANGTGCACT 120
 TTTNTTTTGG NAGTTCAGAN TCCAN 145

SEQ ID NO:6656

SEQUENCE LENGTH:351

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07850

SEQUENCE DESCRIPTION:

GATCCTCTGG TTTNTTCCAT TGGGGCGGAG TCGGGGGTGG AGGGAGCTGG CCACAACCCA 60
 CTGCTCTGAT GGGTGGTTT TCCAAGNATG CTGAATGTAA TGCCTGGTCA ATGTGGAAGC 120
 CCATNAGGTT GCCCAGGGAA GCCTCCAAAA GCTGGGATGC TTGAGGGTAT CCAAGTTGAA 180
 AAAGACAAAA TCTGACCATC AGCCAGTGAC AGTCCTGGCA AATNAAGGTG GGGCGGGGCA 240
 GTNAGGGGTG GNGANGGTG AATNNTTCAT TATTCCACCC NGCGGTTTNC TGGGGTGAGG 300
 GGAAGANTC GNTGCTTGCT TTGGGCACTG ANGGTTTTTC TGTNNGGGGA N 351

SEQ ID NO:6657

SEQUENCE LENGTH:346

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07852

SEQUENCE DESCRIPTION:

GATCATGTGA AGAAAACAGT GACCATTGAA AATNACCCTC ATCTGCCACC ACCTCCCATG 60
 TGTTCACTTC ACCCATGCAG GCATGCTGAG GTGATGAAGA AAATCATTGA GACTGTTGCA 120
 GAAGGAGGGG GAGAACTTGG AGTTCATATG TATCTNCTTA TTTCTTGAA ATTTGTACAA 180
 GCTGTCATTC CAACAATAGA ATATGACTAC ACAAGACACT TCACAATGTA ATGAAGNGAG 240
 CATAAANCT ATCCTAATTA TTGGTTCTGA TTTTNAAGA ATTAACCCAT AGATGTGNCC 300
 ATTGNCATA TTCATCANTA TATACAGTTT CTCTANTAGG GGAANN 346

SEQ ID NO:6658

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07853

SEQUENCE DESCRIPTION:

5 GATCCACACG CTGGCCCAGG AGTTCGACAT TNACCGCAAG GTGGGCGNAG AGCCTGTNCC 60
 TGTAACGAGG GATNATTCCA GCAATGGCTT TCCCCGCACT CAGCATGGTT CCCCAGTAAC 120
 CTCCTGTGTA CCAGCTCCTC GGGGGAGCTG TGGAGGATGG TGGCATCGG TGGACAGCCC 180
 CTGGGCTTTA ATGAATGTGG CATCGTGGCA CAGATTGCAG GTCCCCTGGC TGCCGCTGAN 240
 ATCTCTGCCT ACTACATCAG CACCTTCAAC TTNNNGNNNG GCCTGGTTGC CCGAGGACGG 300
 10 TATCGGCAGC GTCATCGNGG GTCCTCCAGN GGNGGGNAGG AATGN 345

SEQ ID NO:6659

SEQUENCE LENGTH:47

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS07854

SEQUENCE DESCRIPTION:

GATCTTTGTT AAAAAAATAA AAAGTACTAA CATTACAGAC ATGTAAA 47

20 SEQ ID NO:6660

SEQUENCE LENGTH:35

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

25 CLONE:HUMGS07855

SEQUENCE DESCRIPTION:

GATCCTGCCA CTGCAGAGTG AAACCCTGTC TCAAA 35

30 SEQ ID NO:6661

SEQUENCE LENGTH:279

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07856

SEQUENCE DESCRIPTION:

35 GATCCCCCG AAGTCTTCAC AGGCACTGCA TCGGGTTGTC TGGCGCCCTT TTCCTCCAGC 60
 CTAAACTGAC ATCATCCTAT GGACTGAGCC GGCCACTCTC TGGCCGAAGT GGCCGCAGTG 120
 TTGCCCCCGA GCTGCCCCCA CCCNCTACA GGGTCCCTCA GNTTATAGGT GCCCAGGCTG 180
 AGGTGAAGNG GCCTGGGGGN CCTGCNTTCC GGGCGCTCCT GGACCNTGGG GCAAACCTGT 240
 40 GACCNTTTN TACTGGANTA GAAATNNGTT TTATNATTN 279

SEQ ID NO:6662

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS07857

SEQUENCE DESCRIPTION:

50 GATCATAATA TAATTGGTTA CAAAGTGGAA GTGCCAGCTG GCTTAAGTAC CCAAAGAANA 60
 GAATGCAGCA GCCTAACTTA GTGTACCAT ATGTTACTGA ATTTGAAACT GACCTTTTTT 120
 CCCACCCTAC TTCACACACC TAAACTCTT TTTTGTGAG ACCAAAGNGC AAA 173

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SEQ ID NO:6663
 SEQUENCE LENGTH:295
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07858
 SEQUENCE DESCRIPTION:
 GATCCAGTTG TCATCCACCA CAGACATCTC ACATCAGATA CAGACAGTTC CAAGATTGAC 60
 AACAGAGAAC AACCTGCTGG AAAGACCTGG GCAGAAATGG AGAGCCCTGC GGAACCATG 120
 CTACATTTTC ATCTAAAGAG AGAATGCACA TCTGATGAGA CTGAAAGTNC TTTNTTGT 180
 TAGATTGTAG AATGGTATTG ANTGGTCTG TGGAAAATTG CATTGCTTTT ATTTCTTTGT 240
 GTAATCAAGT TTAAGTAATA GGGGATATAT AATCNTAAGG CNTTTTAGGG TGGCN 295

SEQ ID NO:6664
 SEQUENCE LENGTH:285
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07859
 SEQUENCE DESCRIPTION:
 GATCATGGTC AGANTGGAGT ACCTGGGGGC TGTCCATGCC CCCCTTTGGA CCTAATCCTA 60
 CCCGTCCCCG CCAGCCCTC TCCACACCNT TNCTCCCCAA GTACCCGCC ACCGTTTCCA 120
 TGGTCGAAGG TCAGGGCGAG AAGAACGTGA CCTTCTGGGG GAGACCGCTG CCACGGGTGT 180
 AGGAGCTACA AGGGCAGAAG CTGGTGGTGG AGGNGNNACG ACCATNTNTA CACGTGCCTG 240
 CTTGCAAAGA CCCTGAGGAA GAGGAACTCT AACANTTNTT TCCTN 285

SEQ ID NO:6665
 SEQUENCE LENGTH:47
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07860
 SEQUENCE DESCRIPTION:
 GATCCTGTTA GTCACGTGG TTCAACAAAT AAAACTGTTT GTGCAA 47

SEQ ID NO:6666
 SEQUENCE LENGTH:75
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07861
 SEQUENCE DESCRIPTION:
 GATCTCTAGA GCTGTCTTGT CGCCGCCAG GATTGACCTG TGTGTAAGTC CCAATAAACT 60
 CACCTACTCA TCAA 75

SEQ ID NO:6667
 SEQUENCE LENGTH:286
 SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

5 TOPOLOGY:linear
 CLONE:HUMGS07862
 SEQUENCE DESCRIPTION:
 GATCCTGCCT GGNCAACCCT TTCCTGTCCC AGGCCCTACA TTCAGCAGAA ACGCACTTTG 60
 GACTTTNTCC TTTANATAAA AGAAAGACAT CCCAGGAGAA GGACAAACCA GAGGAGTGAA 120
 CCAACAAAGA GTACCTAGGA ATGGAAGTTG AGCCCTGGAA TGGGGCTCCA TGGAGAGGTG 180
 CATAGGACTC GGCAGAAATG GCCTCTCCCC AAAGCCTCTT TTTNAGAGGA GAGGGAAGCC 240
 10 TATTTTNTTA ACTGGTTTGG GATAGGGAAT GGGGTTCNTT TTTNTN 286

15 SEQ ID NO:6668
 SEQUENCE LENGTH:95
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07863
 SEQUENCE DESCRIPTION:
 GATCCCCGCC TGGCCAGNG CAGGCACGGG TGGTCCCNGT TCCACCCCAA GAGAACTCGC 60
 20 GCTCAGTAAA CGGGAACATG CCCCTGCAG CCAAA 95

25 SEQ ID NO:6669
 SEQUENCE LENGTH:213
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07864
 SEQUENCE DESCRIPTION:
 GATCTCCTAA TGATGTATTG TGCCGTGGNA GTACTGTGTG TGAATAGCAG TAGTGGGGGC 60
 AAAAGCAATC TTCTCATTG GAAATGTTGT AAATAATTTT ATTATATAGT GTTTTGGATG 120
 30 TATTTGTTGT AGAAATGGNC CAGTGAATAA AGAGAATCTA AGGNTTTGTA CAATGTGAAA 180
 TAACGTGTTA AATAANTGTC ATTGTCATAG AAA 213

35 SEQ ID NO:6670
 SEQUENCE LENGTH:137
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07865
 SEQUENCE DESCRIPTION:
 40 GATCTAAAGA CCCGGGAGCG AGAAGCCAAG GCCAGGTTCT GGGNGNAGGG CCCAGAGAAG 60
 CAGAACAGCC CAGAGCCCA GGTGCCTGGC CTGGGCTAGA CCTCTGGAAT TGAGATTAAA 120
 CAATTTGGAG TTGAAA 137

45 SEQ ID NO:6671
 SEQUENCE LENGTH:276
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07866
 50 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

5 GATCCGCTAC AGCAACGTGA AGAAGCTGGA AATAAAGCCA AAGTACCCGC ACTGCGAGGA 60
 GAAGATGGTT ATCATCACCA CCAAGAGCGT GTCCAGGTAC CGAGGTCAGG AGCACTGCCT 120
 GCACCCCAAG CTGCAGAGCA CCAAGCGCTT CATCAAGTGG TACAACGCCT GGAACGAGAA 180
 GCGCAGGNTC TACGAAGAAT AGGGTGAAAA ACCTCAGAAG GGAAGACTCC AAACCAGTTG 240
 GGAGGCNTTG TGCAAAAGGN CTTTTCAGN TTAAAA 276

10 SEQ ID NO:6672
 SEQUENCE LENGTH:348
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07868
 SEQUENCE DESCRIPTION:
 15 GATCCTGGAC CAGTCAGAAG CCGAGAAAGC TCGCAAGGAA CTTTGGAGC TGGAGGCTCT 60
 GAGCATGGCC GTGGAGAGCA CCGGGACTGC CAAGGCGGAG GCCGAGTCCC GTGCGGAGGC 120
 AGCCCGGATT NAGGGAGAAG GGTCCGTGCT GCAGGCCAAG CTAAGACAC AGGCCTTGGC 180
 CATTGAAACG GAGGCTGAGC TCCAGAGGGT CCAGAAGGTC CGAGAGCTGG AACTGGTCTA 240
 TGCCCGGGCC CAGCTGGAGC TGGAGGTGAG CAAGGCTCAG CAGCTGGCTG AGGTGGAGGT 300
 20 GAAGNAGTTC AAGCAGGTTG CCAGAGGCCA TAGGCCCAGC ACCNTNAN 348

25 SEQ ID NO:6673
 SEQUENCE LENGTH:325
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07869
 SEQUENCE DESCRIPTION:
 30 GATCTGAGGC CAGCTCCCGT TCCCTGAGCT GCCTCCTGCT TCTCCCTCAA AGTCTCCTAC 60
 CCTTCTCTTC CTCTTTNAAG CCCTCTCTTC CTCGCTTTCC TTCCTACCTA GCTCCTTGGT 120
 GGTGAGCTTN TTGTGCCTTA ATNCTGTGAC CCAGGCCCTT ACACCANTTT NCACCTTCCT 180
 GTNCGGAGNA CAAGGNACT AGCTGCCCCA GGAAGTTGTG TGATTTTAAA TCACTTCTGT 240
 CTTTGTGGA AAGTGATTT GTGCATAAAT AAAAGTCTGN GGATTTGTTA AAAANAAGA 300
 35 AAAANACNNA AAAANAACAN TNAAN 325

40 SEQ ID NO:6674
 SEQUENCE LENGTH:348
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07870
 SEQUENCE DESCRIPTION:
 45 GATCAATGGA ATAGAATTTT ATTGGAAATG TAAACACTT TCCCAACAAT GGTCATGACT 60
 TTCTNCTGTT TTTGAGAAGA GTTTCATATG CTGGACCACA TTTNAGCTTT TATTGTTTTT 120
 TTTTNCCTAT TGTCAAAAA GTTAAGCAAC AAGTGGCCAC ACTTTTACGT GNCTACAACC 180
 TGGAGTTCTG CAAAGAAGGT AATATTTNCT TGGNCTTTGC CTAANGTTAT CTCCCCATTG 240
 TATGGTTACA TTTTATTTTG GCCTATGGGG CTTTCTAATA CGTTTTGGTA AAGAGGCGAG 300
 50 TNTAANGNAA TTNCTTGTC AATTTCACCT CAAAGGTAAT TTCNTGCN 348

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SEQ ID NO:6675

SEQUENCE LENGTH:196

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07871

SEQUENCE DESCRIPTION:

GATCAACTCT CTAATGGTGC TAATGGCAAT CTAGCTAATG TGCAAATTTA GGAAGTCTTC 60
 AGTACTAAGT ACATAATTTT CAAATAATAT TTTTNAATTG TCTACTTTGG ATGTTAGCTA 120
 TGTCTTGTGN GTATAGGATT CCAATTTTAA GCACTATTTT GATGCAAAAN AATGCATAAN 180
 NAATTTANTT TATAAA 196

SEQ ID NO:6676

SEQUENCE LENGTH:214

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07872

SEQUENCE DESCRIPTION:

GATCATAAAA ATGTGCGCCA ACAACGGCAG GCGCCATCTA AAGCAGCTTC TAAACAGAGA 60
 GAGATGCTCA TNGAAGATGT NGGCAGTGAG GNAGAACAAG AAGAGGAGGA TGAGGCACCA 120
 TTCCAGGAGA AAGATTCCGG CAGCGATGAA GATTCCTAA TGGACGATGA TGACGNTNGT 180
 NACTATGGCA GTTCGAAAAAN' GAAAAACANA TNNN 214

SEQ ID NO:6677

SEQUENCE LENGTH:138

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07875

SEQUENCE DESCRIPTION:

GATCCCCCAA GCCTGACGNG CCGNTNACCA TCCCCTCTGC CCTGCAGAGN CAGCCGCCAA 60
 GGNTGACCTC AGCTTCGGAG CCACCTCTGG ATGAACTGCC CCCAGCACCC GCCCCATTAA 120
 AGACCCGGAA GCCTGAAA 138

SEQ ID NO:6678

SEQUENCE LENGTH:208

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07876

SEQUENCE DESCRIPTION:

GATCCATCGC AGAGTCCTAA AGAAGAACCC ACTGAAAAAN TTGAGANTCA CAAGCTCCGG 60
 GTGGATAAGG CAGCTGCTGC AGCANGGCAC TACAAGCCAA ATCAGATGAG AAGGCGGCGG 120
 TTGCAGGCAA GANGCCTGTG GTAGGTAAGA NAGGAAAGAN GGCTGCTGTT GGTGTTAAGA 180
 NGCAGAAGAN GCCTCTGGTG GGAAANN 208

SEQ ID NO:6679

SEQUENCE LENGTH:188

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07878
 SEQUENCE DESCRIPTION:
 GATCTCCTGG AGGCTACAGG GAGCAGAGGC GATGCCCACA GCGGCGAGCG TGTTCCTCAGG 60
 AGCAGGTGGG CAGGGGCCTT GCAGACTCTT CATTTCTTCC TTCCCTTATC ACCGTTAGCC 120
 ATTTTATGG GATAGAGATA CAGAGGAGCT CCCATTACAA AGCGCCATTA AACTGTCATC 180
 TGTGCAAA 188

SEQ ID NO:6680
 SEQUENCE LENGTH:35
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07879
 SEQUENCE DESCRIPTION:
 GATCTTGGTG GTAGTAGCAA ATATTCAAAC GCAAA 35

SEQ ID NO:6681
 SEQUENCE LENGTH:297
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07880
 SEQUENCE DESCRIPTION:
 GATCTTCCGT GGTGAGAGTG CCACTGCGGG AGCTCTGTAT GGTGAGGATG TAGGGGTAA 60
 CTTGGTCAGA GCCACTCTAT GAGTTGGACT TCAGTCTTGC CTAGGCGATT TTTTCTACCA 120
 TTTGTNTTTT GAAAGCCCAA GGTGCTGATG TCAAAGTGTA ACAGATATCA GTGTCTCCCC 180
 GTGTCTATC CCTGCCAAGT CTCAGAAGAG GTTGGGCTTC CATGCCTGTA GCTTCTCTGG 240
 TCCCTCACCC CCATGGCCCC AGGCCACAG CGTGGGAACT CACTTCCCT TGTGNTN 297

SEQ ID NO:6682
 SEQUENCE LENGTH:22
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07881
 SEQUENCE DESCRIPTION:
 GATCCAAGAT ATCACTTAGA AA 22

SEQ ID NO:6683
 SEQUENCE LENGTH:292
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07882
 SEQUENCE DESCRIPTION:
 GATCCAGCTG TGGTCCAGA GCAGTGAGCT GGTGAGCTGG GCAGATGGGG GCCAGCACAG 60
 CAGNATCCAC CCAGNCTGAG GCTCAAGCTG CCCTTACCAC CCCATCCCC ACGCAGGACC 120

EP 0 679 716 A1

AACTACCTCC GTCAGCAAGA ACCCAAGCCC ACATCCAAAC CTGCTGTCC CAAACCACTT 180
 ACTTCCCTGT TCACCTNTGN CCNACCNCAG CCCAGAGGAG TTTGNGCCAC CAACTTCAGT 240
 GNCTTTTGT ACCCNAAGNN AGCACAAGAT TGGACCAATA NTTTTGNAG TN 292

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SEQ ID NO:6684
 SEQUENCE LENGTH:247
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07883

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SEQUENCE DESCRIPTION:
 GATCTGCCTT AGCCTCACTG CTGCCCCCTA AGTGGTGGGA TTGCAGGTGT GAGCCACTGT 60
 GCTGGCCTCA AATAGTTTNN ATCAAAGCTA CTTCAATTAG TGGTTAGCAA GGCTTAAAGA 120
 CTAATTCAGT GCTTTATTTT NACACTNGNT CGCAACATGT TGCTTTTTTN CCTGTGTCCT 180
 ATGGGACCTA GTCATCTGTA TGTNAGATTC ACAGAGGAAT TAAAGAGATA AACATGCATG 240
 TNCTAAA 247

15

SEQ ID NO:6685
 SEQUENCE LENGTH:37
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07884
 SEQUENCE DESCRIPTION:
 GATCCACAAC TACACACATC ATAACCACAC CATCAAA 37

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SEQ ID NO:6686
 SEQUENCE LENGTH:81
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07885
 SEQUENCE DESCRIPTION:
 GATCTCTGAG GAGTTCCTG GGGACAACCTG AGCAGCCTCT GGAGAGGGGC CATTAAATAAA 60
 GCTCAACATC ATTGNACNAA A 81

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SEQ ID NO:6687
 SEQUENCE LENGTH:237
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07886
 SEQUENCE DESCRIPTION:
 GATCAGCCTT TCCAGGTGCT TTAGTCGTTT GCCTAGACAC GTTCCTGTCG CCTCTGACAG 60
 CTGCAGTTGC TCTGCCTGCT TAGACATGGT GGACATTINA GCAGCCTGAC CTGTGGGGAG 120
 GGGGTCTCTC CCGAAGAGTT TCTGTTTTTA CTCAAATAAA TGTTATTCTC AGATGCTTGA 180
 TGCACGTGTT GAAATGTGCT CNATTTAATC ATGCAGATAA ACCATTAAAA TGTCAAA 237

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SEQ ID NO:6688

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EP 0 679 716 A1

SEQUENCE LENGTH:93
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
5 CLONE:HUMGS07887
SEQUENCE DESCRIPTION:
GATCTCTTGA GCCCAGGAAG CCGAAGCCAC AGTGAGCTGT GATTGCACCA CTGCACTCCA 60
GTCTGGGTGA CAAGAGTGAG ACCCATCTC AAA 93

10 SEQ ID NO:6689
SEQUENCE LENGTH:62
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
15 CLONE:HUMGS07889
SEQUENCE DESCRIPTION:
GATCGCAGCA TTGCACTCCA NCCTGGGCAA CAAGAGTGAA ACTCTGNCTC AGAAAAGAGA 60
AA 62

20 SEQ ID NO:6690
SEQUENCE LENGTH:221
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS07890
SEQUENCE DESCRIPTION:
GATCCTCCTG CTTAGCCTC CCAAAGTGCT AGGATTATAG CCACCGCACC TCGTCTTGAA 60
ATAGCCTTTT AAATGTTTAC CTCAGCTTGC CTCACAGTGG GTCTGTTCTT GGTTCCTCAG 120
ATGCATAAAG GAAGACATAT CCCTCCCCTG GGNAGCAAGG CTACAATGGG AGGGAGGGAG 180
30 AACATGGGAG CATGTGAATA AAATGGCATT AANTACTGAA A 221

SEQ ID NO:6691
SEQUENCE LENGTH:210
35 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07891
SEQUENCE DESCRIPTION:
GATCATCAAG AAAAAAGAAA CTAAATGGA CACTTTTAAT GCTGCAAATN AGCTTGTCGT 60
40 GAAGTTACCT GATTGTNTAA TTAGAATGAC TACCACCTCT GTCTGATTCA CCTTCGCTGG 120
ATTCTAAATG TGGTATATTG CAAACTGCAG CTTTCACATT TATGGCATTT GTCTTGTTGA 180
AACATCGTGG TGCACATTG TTTAANCAAA 210

45 SEQ ID NO:6692
SEQUENCE LENGTH:169
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07892
50 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

5 GATCTGTCAT CTTATAGCA CAACAAAACG AAATGATGGA AATNCTCTTN AGCTCACAAC 60
 ATTTGTTTTN CTTTAAAGT AAATGCAAGT ACCAAAGCTC ACTACTGCGG TTTGCCTGTN 120
 CCTGGNCANT GAGGCGGAGC NCTGTTGTTT TGTGGGGCA CCCCCTTNN 169

SEQ ID NO:6693
 SEQUENCE LENGTH:226
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 10 CLONE:HUMGS07893
 SEQUENCE DESCRIPTION:
 GATCTCTGCT TACCGTTCAA GAGGCGTGTG CAGGCCGACA GTCGGTGACC CCATCACTCG 60
 CAGGACCAAG GGGGCGGGGA CTGCTGGCTC ACGCCCCGCT GTGTCCTCCC TCCNTCCCTT 120
 15 CCTTGGGCAG AATGAATTCG ATGCGTATTC TGTGGCCGCC ATCTGCCGAG GGTGGTGGTA 180
 TTCTGTCAAT TACACACGTC GTTCTAATTA AAAAGCGCAT TATAAA 226

SEQ ID NO:6694
 SEQUENCE LENGTH:31
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07894
 SEQUENCE DESCRIPTION:
 25 GATCTCTTAA ATAACTTGC TTCTGGTTAA A 31

SEQ ID NO:6695
 SEQUENCE LENGTH:182
 SEQUENCE TYPE:nucleic acid
 30 TOPOLOGY:linear
 CLONE:HUMGS07895
 SEQUENCE DESCRIPTION:
 GATCTAAATT TGTATTATTA AGCTGCAAGN AAGTTACATT AATTTGGAAT GTGTCATCAC 60
 35 TTGGNCCCAG TATATCAGAT TTTNATTGAG TAAAATGTTA AAAATAATAA GTGTAAGTCA 120
 GATTCTGGAT GTNCGNNGTTT ACAATACATT GCCTGTAATA AAAGCTGATT CTGANGTGCA 180
 AA 182

SEQ ID NO:6696
 40 SEQUENCE LENGTH:203
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07896
 SEQUENCE DESCRIPTION:
 45 GATCCCAACC CTGTGTCCTG CCCTGACCCC TGTTCTCCCA CTCGTGGCAC TGTCAGCCCA 60
 GNCCTCCCTA CCGGCGAGAG TCCAGAGTGG GTACAGGAGC AAGGGGCACT GCTGGGGCCT 120
 GATGGCTGAA GGAGACGCCG GCATCCTCGG GGGCCTGGGG AAGTTGTGTG TTGTGCAGTC 180
 50 AGTAAATCC TCCCACTGCN AAA 203

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5 SEQ ID NO:6697
 SEQUENCE LENGTH:51
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07897
 SEQUENCE DESCRIPTION:
 10 GATCAAAATGA AATAATGTAA CGTGAGNATA CAGTATAAAT GATAAAGGAA A 51

15 SEQ ID NO:6698
 SEQUENCE LENGTH:65
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07898
 SEQUENCE DESCRIPTION:
 20 GATCTCCGGG CAGCCACCAC CTCCTCGGTC TGCCCCNTCA TTA AAAAGTC CTCCTTCCAT 60
 AGAAA 65

25 SEQ ID NO:6699
 SEQUENCE LENGTH:59
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07900
 SEQUENCE DESCRIPTION:
 30 GATCTAGTGC CTAGTGTTC TTTCTGATG TAATAAAAGG TGGTCTGGCA GAACCTAAA 59

35 SEQ ID NO:6700
 SEQUENCE LENGTH:129
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07901
 SEQUENCE DESCRIPTION:
 40 GATCCAGAGC CACAGAAAGA AATGTAGGTG TGAAGTATTA GGCTGCTGTC AGGGNGAGGA 60
 TGGCAGATGG AGGCATCAAG CACAAGGAAA ATGCACAACC TATGCCCTNT TATACANNNN 120
 GTTCATGTN 129

45 SEQ ID NO:6701
 SEQUENCE LENGTH:39
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07902
 SEQUENCE DESCRIPTION:
 50 GATCTTAAGT TTACAATTAA AAATCAGTA CTCAATAAA 39

55 SEQ ID NO:6702
 SEQUENCE LENGTH:56

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07903
 5 SEQUENCE DESCRIPTION:
 GATCCCCCG GGCATGGCCT GGGCTGGTTT TGAATGAAAC GACCTGAACT GTCAAA 56

SEQ ID NO:6703
 10 SEQUENCE LENGTH:31
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07904
 SEQUENCE DESCRIPTION:
 15 GATCCTTTTA AAATAAAAAT NTTGATGCAA A 31

SEQ ID NO:6704
 SEQUENCE LENGTH:108
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07905
 SEQUENCE DESCRIPTION:
 GATCAGGAGC CTTTATTAAA ACAGAAATTT TAGGGAAAAC CACAAAAACC TTCGTAAAGT 60
 25 GAATTGCATT AACATCAAT TATAATAAAA TTTCATATTT TGANTAAA 108

SEQ ID NO:6705
 SEQUENCE LENGTH:29
 30 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07909
 SEQUENCE DESCRIPTION:
 GATCAAAAGG GACAAGCATC AAGCACAAA 29

SEQ ID NO:6706
 SEQUENCE LENGTH:152
 40 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07911
 SEQUENCE DESCRIPTION:
 GATCATGGGC AAAATCTAGC CACAGTCCTG AGAGTCCAGG CTTCTGGGAT GCCCAGCTGG 60
 GTATAAAAGT CCCTACTAAC CCTGNTTCAA ATTCAGAGGT TTCTTTGGTT NAGAATGCCT 120
 45 CAATGAGATT TTGATACATC CAAGAGCGCA AA 152

SEQ ID NO:6707
 SEQUENCE LENGTH:129
 50 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS07912

SEQUENCE DESCRIPTION:

5 GATCTAAAA ATACAAAATG TAGACAATTA CTATTCAGGT GCTTACTTTT AAGCCAATAT 60
GTAGCAGNGN AAAACTAATT TCTCTGTGGC AGTATTTTAA TTTTGGNAT GTCCNNGGNN 120
TTAACTGAN 129

SEQ ID NO:6708

SEQUENCE LENGTH:118

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07913

SEQUENCE DESCRIPTION:

15 GATCCGCACT GGGGGTCTG CAGGGTGGG NCAACCACAG CCTGTCACAA GTCACCGTC 60
AGCCTCTNTG GNGCCATTG GCAGCATGAA GTCCCTATCA CCCACATAAT AATGNGNN 118

SEQ ID NO:6709

SEQUENCE LENGTH:149

20 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07915

SEQUENCE DESCRIPTION:

25 GATCTTTCCC AGTATCTNNA GTGGGTGACC ACACTTGTC GTGGGAGNCT NTGAAACTAC 60
CCTATCGNCT CCTTAAGGGC GGGATGAACT GCTTTGTGAC TTGGAAAGGT ACGCTGCTGG 120
CCAGCATTGG AGANGAAGCT NCTGAGCAN 149

SEQ ID NO:6710

30 SEQUENCE LENGTH:236

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07916

SEQUENCE DESCRIPTION:

35 GATCAGGTGG TCAAGGTGCT GCACGATGCC CAGCAGCAGT GCCGAGACGG CCACCCGGTT 60
GTGTCAACA TCCTCATTGG GAGGACGGAC TTCCGCGATG GCTCCATTGC TGTATAGGGC 120
CTTGTGGGTC AGNACGCTTN GCTGCCTTCC TACCCCTGGA CTGTNTCCAA NTGGTTTNG 180
40 AGTCTCATNA TTGCTTACCC CTGGNGCTNA TCCCATGAGG NCCCANTGNA CTTCAN 236

SEQ ID NO:6711

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

45 CLONE:HUMGS07917

SEQUENCE DESCRIPTION:

GATCCGCAAA GAGGAAGACC GCCTCTTCAT TCTCAGGACC NTCACCTATT GATGGGAATN 60
TGTGCGGGCT CAGGGTTCTT NNACACACTA GNTGTGGGAA GCCATAGGAG CCTCCAGATG 120
50 GGGGCTGGCC TCTTTGCC AGCNAGNGGG CAGGGCCTGT GGGNTGGTGA ATN 173

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EP 0 679 716 A1

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 SEQ ID NO:6712
 SEQUENCE LENGTH:48
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07918
 SEQUENCE DESCRIPTION:
 GATCACGNCA TGGCACTCA GCCTGAGCGN CAGAATGAGA CTATTAAA 48

10
 SEQ ID NO:6713
 SEQUENCE LENGTH:50
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15
 CLONE:HUMGS07919
 SEQUENCE DESCRIPTION:
 GATCGTGTCA CTGAACTCCA GNCTGGGTAA CAGTGCGACT CTGTCTCAA 50

20
 SEQ ID NO:6714
 SEQUENCE LENGTH:156
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07920
 25
 SEQUENCE DESCRIPTION:
 GATCTTTGTT TAAATGCCAA AATGTACTTA AATGAGTTAC TTAGAATGCC ATAAAATTGC 60
 AGTTTATGT ATGTATATAA TCATGCTCAT GTATATTAG TTACGTATAA TGCTTTCTGA 120
 GTGAGTTTTA CTCTAAATC ATTTGGTTAA ATCAAA 156

30
 SEQ ID NO:6715
 SEQUENCE LENGTH:208
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35
 CLONE:HUMGS07921
 SEQUENCE DESCRIPTION:
 GATCGATACC CCCACCCCTC CACTNGCCAG CCTGGGGGTG CCCTGCCTGC CCTCCTGGTA 60
 CTGTTGTCTT CCCCTCGGCC CNNTCACATG TGTATTCAGC AGCCCTATGG CCTTGGCTNT 120
 GGGCCTGATG NGNAGGGGTA GAGGGAAGNT GAGCATAGCA CATTTTCCTA GAGCGNGAAT 180
 40
 TGGGGGNAAG CTGTTATTTT NATATTAN 208

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 SEQ ID NO:6716
 SEQUENCE LENGTH:57
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07922
 SEQUENCE DESCRIPTION:
 GATCACGACA CTACACTCCA GCCTGGGGGA CCGGGCAAGA CTCTGTCTGT CTTAAA 57

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EP 0 679 716 A1

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SEQ ID NO:6717
SEQUENCE LENGTH:333
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07923
SEQUENCE DESCRIPTION:
GATCTTGAAA AAAAAAGCAA AAGCTTAAAT ATTTGATACA AGTTTACTTA GCTACAACAT 60
ACTTTACATT GTTGCCTTTA GTTATCTCAC AGGCACTGAC ATTTTATATT TAGAAAAATAC 120
TTTAAATCTT NCTAATCTTT TTTTGAAAT ATTAGTGTCC ATTCTGTATG ACTCGCTAAC 180
CTACTTTGCA AGGCTTTGGG CAACATTTTA GTCATTAAC TTCAAGATGA TGTGTCATCT 240
GTATAGGTCA AAGAATGGGN CTTCTGAACT GAGGAATTTT GCTGTTGACA GCCAAAGTNT 300
AGTGTACAAG GTTTGATGNA ACTNGNTTAT GTN 333

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SEQ ID NO:6718
SEQUENCE LENGTH:57
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07925
SEQUENCE DESCRIPTION:
GATCTTGGAC TTCCAGCCT CCAGAATTGT GAGAAATAAA TTTCTGTTGT TGATAAA 57

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SEQ ID NO:6719
SEQUENCE LENGTH:63
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07927
SEQUENCE DESCRIPTION:
GATCTTTGCT TAATAAGANG AAAAAAGAGG CAAGGAGAAA ACGAAGAGAA AGAAATTTTG 60
AAA 63

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SEQ ID NO:6720
SEQUENCE LENGTH:22
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07928
SEQUENCE DESCRIPTION:
GATCCNGTGT TCTTCTGTGA AA 22

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SEQ ID NO:6721
SEQUENCE LENGTH:186
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07932
SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

5 GATCCGGCTT GGACAGGCAC CTGAGATGGT GCCAAAGTGC AGCTGACTCT TCCCACGACA 60
GCCCTGCCCT TCCCATGAGG CAGGCTCTTC AGTGAGTGT TGAACGTAAT TATGTAGTTT 120
TCTGTTTAAT TGA AAAAGNG AGCTATGCCT TTTTTCCTT TTGGANGTAA AGCAGCTAAA 180
ANCAAA 186

10 SEQ ID NO:6722
SEQUENCE LENGTH:45
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07933
SEQUENCE DESCRIPTION:
15 GATCCAGGTG TTTGTTTCTA ACTTCTGTAA TACATACAAT GCAAA 45

20 SEQ ID NO:6723
SEQUENCE LENGTH:173
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07934
SEQUENCE DESCRIPTION:
25 GATCATGGTG ATTTNACTGC TGCTTGTGGN TATCGTGGTT GTTGCACTCT GGCCGACCAA 60
CTGATGGCAG TAAAGAGACC ACCAGCAGTG ACACCTGCCA ATNACAGATG CAAGCCCAAC 120
ACCCTNTTGG TACGCAAAAC CTNATCTCAA TAAATTCCCC CAAAGCTCTG AAA 173

30 SEQ ID NO:6724
SEQUENCE LENGTH:61
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07935
SEQUENCE DESCRIPTION:
35 GATCCCAAGC CCTGCATTGT AAGCCACAG TAGGCACTCA ATAAATGCTT GTTAAAGGAA 60
A 61

40 SEQ ID NO:6725
SEQUENCE LENGTH:269
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07936
SEQUENCE DESCRIPTION:
45 GATCCTGCAG AAGAATAACC TGAATCTGCT CAGAGACCTA AGCTGTGCAC ATTGCCATA 60
GCCTCAGGAA CAGTCCAGAC TGGGGCGGTG TGGTCATATT ACACAGNNAG GAGGGTGATT 120
CAGAGTTCAT GAATATCATT GCCAATNAGA TTGGGTCAGA GGAGACCTC CTGTTCTTAA 180
CTGTGCGCGA TGAGNAAGNT GGTGGNCTNT TTTACTGGC AGGGCCACNA NCGCTCATGT 240
GGAGACCCNG GGGCCAGGG TGGCTGATN 269

50 SEQ ID NO:6726

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EP 0 679 716 A1

SEQUENCE LENGTH:229
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07937
 SEQUENCE DESCRIPTION:
 GATCCCTCAT CCCTGAAATT GAGTTTATGT AGTCATTTTA CTTATTNAT TCATTAGCTA 60
 ACTTTGTCTA TGTATATTN TAGATATTGA TTAGTGTAAT CGATTATAAA GGATATTTAT 120
 CAAATCCAGG GATTGCATTT NGAAATTATA ATTATTTCTT TTGCTGAAAGT ATTCATTGTA 180
 AACATACAA AATAAACATA TTTNAANACA TTTGCATTTN ACCACCAAA 229

SEQ ID NO:6727
 SEQUENCE LENGTH:50
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07938
 SEQUENCE DESCRIPTION:
 GATCCATGAA CTCTGACCCC TCCCCAGTAA AGGCTTCTGT AGAGAGCAAA 50

SEQ ID NO:6728
 SEQUENCE LENGTH:122
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07939
 SEQUENCE DESCRIPTION:
 GATCCCCGCA CCAGCCTCTC CTCTCTTTCC CTGCAACCTG TTACCTCACA CAGCTGCTCA 60
 GGAACAGCAG AAATAGTAAA ACTCTTCCAT ATATGTGGCA CCTCAGCCGC CCCTCCNTTA 120
 AA 122

SEQ ID NO:6729
 SEQUENCE LENGTH:213
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07941
 SEQUENCE DESCRIPTION:
 GATCAGGAAG TTCTTAGATT CTTGGATGAC AGATGCATGT TGATGCCCTA TGGAGATGTC 60
 CTINTGTTTT GAGGTCACTG AGGTAGGAAG ACCTGTCTAC TCTTGGTTTC ACCACTAGAA 120
 CAGTCTTGGG CTGGATGGGT TATAGAGCTG AGCGGCTGTA ATGGTTCTNT TTTTACATTA 180
 ACAAAAACAA TAAAAACAC CAAAAACAAC AAA 213

SEQ ID NO:6730
 SEQUENCE LENGTH:85
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07942
 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

GATCCCAGGN ATTCAAGGTT AGAGTGAGCT ATGATTGCAC CACTGTACTC CAGCCTGGGC 60
 ANCAGAGANN GACCTGTCTC ACAA 85

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SEQ ID NO:6731
 SEQUENCE LENGTH:313
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS07944

SEQUENCE DESCRIPTION:

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GATCCAGCTG GAATTGAGG AACTGGAAC AATGAGGTCA GCTTCCAAAA CAAAACAAAA 60
 CAAAACTTTA AACATCTGTG TAAGTACAGG GTGGAATATA ATGAATTCCA GGTACCCATC 120
 ACTTAGGTTC CACAATTAGC CATGGCCAAT CTTGTGTGAT AGGTATGTCC CTACCTATTT 180
 CTCCCCTGCC CTNATTGTGA AGTAAATATT TTCCTAAAAG GTCTTTTAA AATTAAGTTA 240
 TAACCATGCC ATCATTGTAC AAANNTAATG CTATTANNTT CCTTANTNTN ATCAAATNTT 300
 CNNGNCTNAN NTN 313

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SEQ ID NO:6732
 SEQUENCE LENGTH:73
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS07945

SEQUENCE DESCRIPTION:

GATCACTGA ACCTAAGAGT TTGAGGCTGC GCACTGCACT CTGGCCTGAC AGAGCGAGTC 60
 ACTGTCTCTT AAA 73

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SEQ ID NO:6733
 SEQUENCE LENGTH:257
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS07946

SEQUENCE DESCRIPTION:

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GATCTGAATT CTAATGCTGG CTTGTGCCTC TAACTATGTG GTTTGGTCAT TATAGATAAT 60
 GCCCAGTTTA ANTTTCCCA TCTTTGAAAA GGAGATAATA TGTNATCTCT TGGGAGTAGT 120
 TCTGAATATG AAGTGTGTTG ATACAAGAAG CACTGACAAT GTTCTGTNN NNNAAATTA 180
 AAACCTGGCCT GGTTCGCCTT TTAAATCAAG NNGCCTTAAC AGGTAAAAAN TTGGAATTAG 240
 NCCTAAAAAN AAAAACN 257

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SEQ ID NO:6734
 SEQUENCE LENGTH:148
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS07948

SEQUENCE DESCRIPTION:

GATCCTTTAG CCTCTGGNA GTGCCCCAGT TGTACCCCT ACACACCCCT CTTGGCATTG 60
 AGTGCCAGTC CTCTGCCAGG CTNTGTGTTA CAAGTTGGGG NGGCGGCAA AGTACCGAAT 120

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TAAAGATGTC AGTTCTAAAG GAGGNAAA

148

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SEQ ID NO:6735
 SEQUENCE LENGTH:289
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07949

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SEQUENCE DESCRIPTION:

GATCATATCC AATTCATATG GAAGTCCCGG GTNTGTCTTC CTTATCATCG GGGTGGCAGC 60
 TGGTTCTCAA TGTGCCAGCA GGGCCTCAGT ACCTGAGCCT NAATCAAGCC TTATCCACCA 120
 AATACACAGG GAAGGGTGAT GCAGGGAAGG GTGACATCAG GAGTCAGGGC ATGGNNTNGN 180
 NAGATGANTA CTTTGCTGGA TGAAGCAGGC TGCAGGGCAT TCCAGCCAAG GGCACAGCAG 240
 GGGACAGTGC CGGGAGGTNT NGGGTAAGGG AGGGCAGTNA CNTNAGAAA 289

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SEQ ID NO:6736
 SEQUENCE LENGTH:114
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07951

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SEQUENCE DESCRIPTION:

GATCAGGCTT TAAATACTC ATTTACAAGT TTTCTATCCT CCTTCAGTGT TAAAGTAGAA 60
 AGTAAAAAGA GTATCTNATA CATGCATGAA ATTAAAGCAT ATACCAAATG CAAA 114

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SEQ ID NO:6737
 SEQUENCE LENGTH:229
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07952

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SEQUENCE DESCRIPTION:

GATCACCCAG CATCCTTCCC CACCCAGCT GTGTATTAT ATAGATGGAA ATATACTTTA 60
 TATTTTGTAT CATCGTGCCT ATAGCCGCTG CCACCGTGTA TAAATCCTGG TGTATGCTCC 120
 TTATCCTGGA CATGAATGTA TTGTACACTG ACGCGTNCCC ACTCCTGTAC AGCTGCTTTG 180
 TTTCTTTGCA ATGCATTGTA TGGCTTTATA AATGATAAAG TTAAAGAAA 229

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SEQ ID NO:6738
 SEQUENCE LENGTH:295
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07953

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SEQUENCE DESCRIPTION:

GATCTCTTGT NCTTTAATTT TAGGGTCTTG TTCCAGGACT CAAATCAGTA ACTTGGTGAT 60
 TACAAGGTGC TGAATGTGTT GGTAACCATA TCGCAATACA CCTCAAGGAA AAGGTTTCAGA 120
 TTTTNAATTT TAAAATATTT NCATTTTTC CTTGAATTN ATATCCGTTT GTCCACTCGT 180
 ACATGCCTAG CCTACAGAAG GGGATATATA TNATGANATG GTCATTTTCC TGAAGNGNAT 240
 ATTTTGCTTG AAATGCAAAG GNCTGAAAGN GNTTGTAGG TTGTTGTTT TTTCN 295

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EP 0 679 716 A1

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SEQ ID NO:6739
SEQUENCE LENGTH:85
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07954
SEQUENCE DESCRIPTION:
GATCATAACA CTA CTCCAGT CTGGGTGATG AAGTGAGACC TGTCTACAAA AAAAAGAAGC 60
CCAAACTAAA CCTAAAGCAA GCAAA 85

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SEQ ID NO:6740
SEQUENCE LENGTH:145
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07955
SEQUENCE DESCRIPTION:
GATCAGGCTG GTCTCAAACCT CCTCGGTTCA AGCAATTCGC CCACCTTGGC CTCCCAAAGT 60
GCTGGGATTA CAGGGGAGCC ACTGCACTGG CCTTCATTGT CTTTGTGCTG CACAACCTAA 120
AAAACCAGTG ACCCTGTATT GGAAG 145

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SEQ ID NO:6741
SEQUENCE LENGTH:81
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07956
SEQUENCE DESCRIPTION:
GATCCTAACA TGTATTGATT GTAAAAGTGA AATATGTAAT AGCCACACTT CACTGTTTTA 60
AAATAAAAGT GCAATTTCAA A 81

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SEQ ID NO:6742
SEQUENCE LENGTH:50
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07957
SEQUENCE DESCRIPTION:
GATCTTGTCT CCAATTAAC CGAGGCTTTC ACCGATAAAA AAAAAACAAA 50

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SEQ ID NO:6743
SEQUENCE LENGTH:115
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07958
SEQUENCE DESCRIPTION:
GATCTGATGA AGTATATNTT TAATTGCCAT TTTGTCCTTT GATTATATTG GGAAGTTGAC 60
TAAACTTGAA AAATGTTTTT AAAACTGTGA ATAAATGGAA GCTACTTTNA CTAAG 115

EP 0 679 716 A1

5 SEQ ID NO:6744
 SEQUENCE LENGTH:175
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07959
 SEQUENCE DESCRIPTION:
 GATCGGGCTA CTACAAAGTT CTGGGAAAGG GAAAGCTCCC AAAGCAGCCT GTCATCGTGA 60
 10 AGGCCAAATT CTTCAGCAGA NGAGCTGNGG AGAGATTAAG AGTGTGGGG GGGCCTGTGT 120
 CTTGGTGGGT TTAAGGCNCA TTGGNGGGNA NTTCAATTNAA TTNTTACTTC TTTN 175

15 SEQ ID NO:6745
 SEQUENCE LENGTH:61
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07960
 SEQUENCE DESCRIPTION:
 20 GATCAAAAGT AACACGTNNA GACTCTGAGC TTAATCCATT AAATGGATTT NAGCATCTAA 60
 A 61

25 SEQ ID NO:6746
 SEQUENCE LENGTH:118
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07961
 SEQUENCE DESCRIPTION:
 30 GATCATNAAC TAACTGGTT ATNTGGAAGT CTAGGTTCTG GTCCTGCTAC CCCTTTAACT 60
 TGAGTNATTT ACAAGTCTGT TTAAGGTTT CTAGACATAC ACTGGTTTGN TTTTAA 118

35 SEQ ID NO:6747
 SEQUENCE LENGTH:140
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07963
 SEQUENCE DESCRIPTION:
 40 GATCGCCCAA CGTTTTTGGN TTATAGAAAT ATTATTCCT GCTTCTTTC TTTGGGTCTT 60
 TTGAATTTCT TTGGTTTCGT NTTAAGAAG TAACCCANCA TTTCTACAA CACTAAATAA 120
 AATGGTACTT ACCTTTCAA 140

45 SEQ ID NO:6748
 SEQUENCE LENGTH:68
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07964
 SEQUENCE DESCRIPTION:
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EP 0 679 716 A1

GATCCCCGTT CCACCCCAAG AGAACTNGCG CTCAGTAAAC GGGAACATGC CCACTGCAGA 60
CACGTAAA 68

5 SEQ ID NO:6749
SEQUENCE LENGTH:79
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07965
10 SEQUENCE DESCRIPTION:
GATCCCCCT GCCTATCAGC TTCTCCTGTG GNGCCTGTTC CTCACTGGAA ATTGGCCTCT 60
GTGTGTGTGT GTGTGTGTN 79

15 SEQ ID NO:6750
SEQUENCE LENGTH:28
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07966
20 SEQUENCE DESCRIPTION:
GATCGAAAGG GAGGCAGGTT TGCCTAAA 28

SEQ ID NO:6751
25 SEQUENCE LENGTH:294
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07967
SEQUENCE DESCRIPTION:
30 GATCTCAAGC CAGCCCCNTC CAGCTCATGA CACTGTTTGG CCTTTCTTGG GGAGAAGGCG 60
GGGTATTCCC ACTCACCAGC CCTGGCTGTC CCATGGGGAA ACCCTGGAGC CATCCCTTNG 120
GAGCCAACAA GACCGCCCCA GGGCTATAGC AGTAAGAACT TTAAGCTCA GGAGGGTGAC 180
GNCCAGATNC GNCTGCTGGG AAGAGCTCCC CTNACAGCT GCAGCTGAAT GCATTAGGGC 240
35 TACCGNAGGN CCCGGACTTA NCAAACTTG NCNACAANNG TTCCTAGGGN TTN 294

SEQ ID NO:6752
SEQUENCE LENGTH:160
SEQUENCE TYPE:nucleic acid
40 TOPOLOGY:linear
CLONE:HUMGS07969
SEQUENCE DESCRIPTION:
GATCCCCAAG TGTGAACAGT CTGAAAAGGG CTTTTGCTT GCTGCAGGGA CCTGAGAATT 60
45 GAGAAACAGC TGTGAAACTT TGAACCCTGG CTGGAAGGTC CTGGTATTGA CTGTGGCANG 120
AAAACGGGGG AGCTAATCTG TAATCTGCTT AGACGCCAAA 160

SEQ ID NO:6753
SEQUENCE LENGTH:98
50 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

5 TOPOLOGY:linear
 CLONE:HUMGS07971
 SEQUENCE DESCRIPTION:
 GATCAGATTC ACGGACCCAG AGCTTTTCCA TGTGTTTATA TTGTAAATAT TTTTNATTTC 60
 ATCAAATTAT TTATTCATTA AAAGAAATTT TTGTGAAA 98

10 SEQ ID NO:6754
 SEQUENCE LENGTH:35
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07973
 SEQUENCE DESCRIPTION:
 15 GATCCTGTCT CTTAGGAAAA AAGAAAACCT CTAAC 35

20 SEQ ID NO:6755
 SEQUENCE LENGTH:64
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07974
 SEQUENCE DESCRIPTION:
 25 GATCAAAGAC CCTGCCTCTA TTTNACCCCG CACTGACTGA ATAAAGCTCC TCTGGCCGTT 60
 NAAA 64

30 SEQ ID NO:6756
 SEQUENCE LENGTH:38
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07975
 SEQUENCE DESCRIPTION:
 35 GATCATCTGA GCTGAAGAAA TAAAAAACA CACACAAA 38

40 SEQ ID NO:6757
 SEQUENCE LENGTH:56
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07976
 SEQUENCE DESCRIPTION:
 GATCATTCAT ACACTATATT TATTTTNC A TTAAANTATT TTAACCTT TGGAAA 56

45 SEQ ID NO:6758
 SEQUENCE LENGTH:105
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07977
 50 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

GATCTTTCAT CAAGATTTTA CCTGCATTTA TGTCTAAGGG TCCCTGATAA GGCTTGAGGC 60
ATGTAACCTA TGTCTAATTA AAAATGATTT GTTGAATGAA NCAAA 105

5 SEQ ID NO:6759
SEQUENCE LENGTH:65
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07981
10 SEQUENCE DESCRIPTION:
GATCTACAGA AGCTTTNATT CTCTTACCTT AACAGTAGTA ATAAAGTTAA CATNTTCACA 60
TCAAA 65

15 SEQ ID NO:6760
SEQUENCE LENGTH:106
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07986
20 SEQUENCE DESCRIPTION:
GATCTGTGTG TTGTGGGAAG AGAATTTTCA ATATGTAAC ACAGAGCTGT AGTGCCATTA 60
GAAACTGTGA ATTTCCACAT AAATCTGAAC ACTTGTCTTT ATTAAT 106

25 SEQ ID NO:6761
SEQUENCE LENGTH:106
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07987
30 SEQUENCE DESCRIPTION:
GATCTCATGG CTTTGATTGN TTTGGGCTGT TCAAAATGTT TATTTGNAAA ACGTATACAT 60
TAATAAACTT AACAAAGCGN TATAAAATAC AGCGNAATCA CCCAAA 106

35 SEQ ID NO:6762
SEQUENCE LENGTH:335
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS07988
40 SEQUENCE DESCRIPTION:
GATCGAAAGG GCCACCATGA ACTTGATACG TCCGTGTGTC CCAGATGCTN NCATTAGTCT 60
ATATGGTTCT CCAAGAACT GAATGAATCC ATTGGAGAAG CGGTGGATAA CTAGCCAGAC 120
AAAATTTGAG ANTACATAAA CAACGCATTG CCACGGAAAC ATACAGAGGN TGCCTTTTCT 180
45 GTGATTGGGT GGGATTTTTT CCCTNTTAT GTGGGATATA GTAGTTACTT GTGCCAAGAN 240
TAATTTTGGC ATNATTTCTN TTANTATCAA CTCTGANGCT AACTTGACT AATCTGAGAT 300
TGTGTNTGNT CCNANTAANC GTGANGTGAA CCTGN 335

50 SEQ ID NO:6763
SEQUENCE LENGTH:169

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07989

SEQUENCE DESCRIPTION:

GATCTTGGAG ACAGGCCATC CATCCTGGAG CCTCATGGAA CAGGATGATG GCACTGAGAA 60
AGCCAATGAC CGAATCTCTT TTCTCTGTAA AAATGTAGAC TGAAGGCCA TGTGTATTTT 120
CCTATGTGCT GTAGCTCTCC TTTGAAATA AATCACAGGC ATCTGGAAA 169

SEQ ID NO:6764

SEQUENCE LENGTH:212

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07990

SEQUENCE DESCRIPTION:

GATCTGGGCT GCAGGAGCTG GGGCCACCCA CAGCCCCCT ACCGACCTGG TGTGGAAGGC 60
ACAGAACACC TGGGGCTGCG GGAACAGCCT GCGTACGGCT CTCATCAAAC TCACTGGGGA 120
AGAAGTGGCC ATGCGGAANT GGTGCGCTCA GTGACTTNTG GTTNAGGACG NCGNGGNTTG 180
AGGATTGAGA NTACNTATT CATTNACCAC NN 212

SEQ ID NO:6765

SEQUENCE LENGTH:141

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07991

SEQUENCE DESCRIPTION:

GATCATACCA CTGCACTCCA GCCTGGGCGA CAGAGAACCT CTCTAAAAA AAAGGAAAAAT 60
NAAAAATAAA TGGGGGCAGA AGCTGAGCCA AGCAGAGGCG GGCAGCTCCT CCTCCAGGGC 120
CCACCTGGCC TGCTGGGGAA A 141

SEQ ID NO:6766

SEQUENCE LENGTH:360

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS07992

SEQUENCE DESCRIPTION:

GATCTGTTAT NAAACGAAAC ACCCCCCGTG TTAATAACTT GGTATGAAAT CTGTTTTTAT 60
GAGCCGGGCC CCCTGTGCCT CTAGTATACT TGTATTGACT CTCATAGTTA CCCTTTTAGT 120
TTTACTGTGT TCTGTGAAAA TTTGTAAATG GTTGNGAATC ACTGTGGGCG TCCATTCTTA 180
TTCAACTAAA TCTCCACAGG TTTTGTGAGC TGGTGTGGAT TAGTTTAACT CTTGTATTCA 240
ACCATTAGTG CTACCACCTT CTCACATTAC AATACANTTN CTGGAAGCAA GTACTGCATT 300
TCCTATGCAA CAAAAAAGG GCAAAANTAC ANANATTGCT AAATGATAAA CAAANACTTN 360

SEQ ID NO:6767

SEQUENCE LENGTH:88

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS07994
 SEQUENCE DESCRIPTION:
 5 GATCAAAAGG GCTATGGGAA GGGCAGACCC CGCCAATGAT TTCTCTTCAC CTGTCTTAAG 60
 ATTAATAAAA AAAGAGTGTC CTGGCAAA 88

SEQ ID NO:6768
 SEQUENCE LENGTH:65
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07995
 SEQUENCE DESCRIPTION:
 15 GATCTTGCCA AGAACCATCA CGTGTTAATT TAGTTGCTTA AACAAAAGGT GTTCTGTTTC 60
 TTAATA 65

SEQ ID NO:6769
 SEQUENCE LENGTH:246
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS07997
 SEQUENCE DESCRIPTION:
 25 GATCAGTGGT GAAATATAGT GATTTTNACC TGTGCTTCCA TTCTGAAGTT CTGGAAAGAA 60
 GTACTGGATG GACTGAAGTC CAGGACAACG TTCCAAAGAA AGGCAGAGTC CAGGTAGGCT 120
 TGGAGGACCA AGCCCTGGAT GAGCACTGGA GGGCAGAGGC CTCAGTGTCC AGCACTGTGC 180
 CCTGCACATG GAAAGCCCT ACGTTTGTGG AATNAATGAN TANTAATAAT ATTTTATNA 240
 GTGAAA 246

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 SEQ ID NO:6770
 SEQUENCE LENGTH:23
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS07998
 SEQUENCE DESCRIPTION:
 GATCCTGTCC CTGCCCTGCC AAA 23

40
 SEQ ID NO:6771
 SEQUENCE LENGTH:46
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS07999
 SEQUENCE DESCRIPTION:
 GATCTGTATA TCCTCCAGTC CAAGATTAAG GAGGCGGACT GTGAAA 46

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 SEQ ID NO:6772
 SEQUENCE LENGTH:108

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EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08000
 SEQUENCE DESCRIPTION:
 GATCCAAACT CATTCTTCT GTAACTCAT GATGACGTAT AACCNITTCC ACCTCACTNG 60
 GAGGTTGGNT CTNCATTCTG AAGGCTTCTA TGTATACATA TTAAATAN 108

SEQ ID NO:6773
 SEQUENCE LENGTH:218
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08001
 SEQUENCE DESCRIPTION:
 GATCAACGAC GACCTTTTTC CTTTNNATAC TAAAGTAAGA AATAAGAATG TNAGCCCCAA 60
 CTGCACTATT TTGCAGACCC CTACCATTTT ACAAAGTGGT CAGAGTGGAA AATCCACCA 120
 GGGCCTGAGC TGTGAGAAAC ATCCTGTCAG GCAGGNCCCA GGNCTNAACC CCTGGCTGCA 180
 CTAANTTCT TCATNATCAN CAGCCTAACA NACCGTCN 218

SEQ ID NO:6774
 SEQUENCE LENGTH:38
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08002
 SEQUENCE DESCRIPTION:
 GATCAACAGT GCAGAGAATC TAGCAAGTAT AAAAGAAA 38

SEQ ID NO:6775
 SEQUENCE LENGTH:252
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08003
 SEQUENCE DESCRIPTION:
 GATCTGCAAC ATGAANCCCG AGGCATGGGT TGGTGTGATG CCAAGGCAAA GTGGTGAGGA 60
 GAAAACAGGA AACGGNCTTT CTCTGAATTG GTAAATGGGA AAGAAGTGAG CAACTTAAGA 120
 TTGTCACANT TAATCACAAG TGTACAGGAT TAGACTGGGT TTATATTTAA CTCTTGCTTC 180
 ATAGGTGTAC CATTAAAGN GTGTTATNTA NTGCTAAGAA NTAAGTGCTT TNATAAAGCT 240
 TATTNNNNAA TN 252

SEQ ID NO:6776
 SEQUENCE LENGTH:34
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08004
 SEQUENCE DESCRIPTION:
 GATCAAAGGG AATAAAGAGA ACTCTTGCT GAAA 34

EP 0 679 716 A1

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SEQ ID NO:6777
SEQUENCE LENGTH:107
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08005
SEQUENCE DESCRIPTION:
GATCATATNC TTTTCTNTT TTGTTCTTA CAATATCAAT CTAATTTAAT ATTCTCTTGT 60
GAGAACACAA ACATNNGTAA ATTGCTCAGT AGTGGCAGCA GTAAGGN 107

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SEQ ID NO:6778
SEQUENCE LENGTH:112
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08006
SEQUENCE DESCRIPTION:
GATCAAGGTT TGTNTGCCCA TTACCTTTCC TCTGCCTGAA AGACGTGTCT CAAGAAAAAT 60
AAATTCTATT TNAGATGCAG GTACTIONATAN TAATTCTAAG ANTTGATATC AN 112

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SEQ ID NO:6779
SEQUENCE LENGTH:140
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08007
SEQUENCE DESCRIPTION:
GATCAGAGNA AACGNCTTTA ACNCCACCAT TAGCACCCAA AGCTAAGATT CTAATTTAAA 60
CTATTCTCTG TTCTTTTCATG GGAAGCAGA TTTGGGTACC ACCCAAGTAT TGAATCACCC 120
ATCAACAACC GCTATGTAAA 140

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SEQ ID NO:6780
SEQUENCE LENGTH:173
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08008
SEQUENCE DESCRIPTION:
GATCAATCTC ATATTCATTC CCTGGGATGT TTAGTTACCA GTTTTCCCAA AGTGTCTCTGG 60
TAGCATCTAC CATATTTNAT CAAATCTGTG ATTCCTTTGA TTATNATATG AACCATTATT 120
TTATGTATCA TTAAGAAAAA ATACTGCCAA TTAAGTCTG TCATGNNAAC AAA 173

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SEQ ID NO:6781
SEQUENCE LENGTH:186
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08009
SEQUENCE DESCRIPTION:

EP 0 679 716 A1

5 GATCCAAGTT GAGATTTTAA TTCTGTGACA TTTTACAAG GGTAAGATGT TACCACTACT 60
TTAATTATTG TTATACACAG CTTTATGATA TCAAAGACTC ATTGCTTAAT TCTAAGACTT 120
TTGAATTTTC ATTTTTTAAA AAGATGTACA AACAGTNTT GNAATAAATT TNAATTCGNA 180
TATAAA 186

10 SEQ ID NO:6782
SEQUENCE LENGTH:102
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08010
SEQUENCE DESCRIPTION:
15 GATCCATTTT TNATATGTCT TGAAGTACTG TGTCTAGTGG GCAAATGTCA TTGTTACCTC 60
TGTGTGTAA GAAAATAAAA ATATTNCTA AAGGTCTGTA AA 102

20 SEQ ID NO:6783
SEQUENCE LENGTH:198
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08011
SEQUENCE DESCRIPTION:
25 GATCAAGTAG ATTTTTTAAA AATCAGAGTC AATTAATTTT AATTGAAAAT TTCTCTTATG 60
TTCCAAGTGT ACAAGAGTAA GATTATGCTC AATATTCCCA GAATAGTTT CAATGTATT 120
ATGAAGTGAT TAATTGGCTC CATATTAGA CTAATAAAAC ATTAAGAATC TTCCATAATT 180
GTTTCCACAA AACTAAA 198

30 SEQ ID NO:6784
SEQUENCE LENGTH:278
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08012
35 SEQUENCE DESCRIPTION:
GATCAGAGTA AACNTGGAG CTAGATTGAT TACAGAGTGC AAGGACTAGG CACCCCGGGA 60
TTGGAGAGCA TCCTTGAAAG CAAGGGTTCT TCATTATTT ATAAAAATT NATTGCCTAC 120
TAGAAGCCCA GCTAAGAGTA TGAAGCAATT AAGACAACAN TCTTTGCTCA AGAATCTCCT 180
AGACAATCCT GATTCAAGCA CCAGAAACAA AGGGNNAAGA AACCTCCAG TCTNGTGNG 240
40 NAAGTGTAC ACATCNTGTC AACAGNTAGG TTNTTTN 278

45 SEQ ID NO:6785
SEQUENCE LENGTH:339
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08013
SEQUENCE DESCRIPTION:
50 GATCCCATGT NAGCTACACA GTGCAGAGGC TCTTGATGGT GGAATAAGCA ATTCCTCCC 60
TCGTGCCAT CTCAGAACCC ATCGGTAGGC AAAGGAAAT ACGCTCAGGT GGTGTGGNN 120

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5 MAGNCTGTNT CAGGCCTACG GAGTCAGCCA GTGGCTAGCG CAAGACCACT CACTCCCTCT 180
 GCNTTCAGGC TTCTTTCAAT TTCATTATCA TCAAGCAGGA ATTATGTCGT AAGTCACTGA 240
 CCCTAACTGC AGACCATGAA GTAAATTATG TAACTAGGTT TTTNNCTTCT CCAGTGGTGA 300
 CCACCNNCNC NCATNCCCCG GTNANAACCT GGGTTTTTN 339

SEQ ID NO:6786
 SEQUENCE LENGTH:80
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08014
 SEQUENCE DESCRIPTION:
 15 GATCGCCTAT TACTGACAGT CTCATTGTAG CTCTAAAAAG CCTAATGTAT CCACTGTGGA 60
 ATAAACTCCA TAGACTCAA 80

SEQ ID NO:6787
 SEQUENCE LENGTH:279
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08015
 SEQUENCE DESCRIPTION:
 25 GATCTTGAAG AGACCGCTGG CAGCACCAGT ATTCCCAAGA GGAAGAAGTC TACACCCAAG 60
 GAGGAAACAG TTAATGACCC TGAGGAGGCA GGCCACAGAA GTGGCTCAA GAAAAAGAGG 120
 AAATTNTCCA AAGAGGAGCC GGTGACAGT GGGCCTNNAG AGGCGGCTNG CAAGAGCAGC 180
 TCCAAGAAGG NAGGAAAAAG TTCCNTTAAG CATTNCCNGG GAGNTTTGAA TTCAAATTG 240
 GGCATTNTTN TTTGGGGTT GGGNNATTCC CTGGCCCN 279

30 SEQ ID NO:6788
 SEQUENCE LENGTH:192
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS08016
 SEQUENCE DESCRIPTION:
 GATCCACTCA CCTCGGCCTC CCAAAGTGCT GGGATTACAG TGCCCGTCCG TGATGATGGG 60
 AACATGNCCG ACGTGCCAG NCACCCCCAG GACCCTCAGG GCCCCAGNCT GGAGTGGCTG 120
 AAGAACTGT GAGCACCTCC ACTGACAGAG GCGGCCCTC CCACGGCTCC CAATAAAAAT 180
 40 GTGAAAACCA AA 192

SEQ ID NO:6789
 SEQUENCE LENGTH:303
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08017
 SEQUENCE DESCRIPTION:
 50 GATCCGTATC GNTTTAAGAA GCGGACGGAG CTGTTTATTG CCGNCGAGGG CATTACACG 60
 GGCCAGTTTG TGTATTGCGG CAAGAAGGCC CAGCTCAACA TTGGCAATGT GCTCCCTGTG 120

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EP 0 679 716 A1

GGCACCATGC CTGAGGGTAC AATCGTGTGC TGCCTNGAGG AGAAGCCTGG AGACCGTGGC 180
AAGCTGGCCC GGGCATCANG GGAATATGCC ACCGTTATCT CCCACAACCC TTNGACCAAG 240
AAGNCCCGTT TTGAANGTTG CCCTNCGGTT TCCAAAGAAN GTTATCTCC TTAANGCCAA 300
5 TCN 303

SEQ ID NO:6790
SEQUENCE LENGTH:267
SEQUENCE TYPE:nucleic acid
10 TOPOLOGY:linear
CLONE:HUMGS08018
SEQUENCE DESCRIPTION:
GATCTAAAA TCAATGAGAA CATAATTTTA ATTGGCAGTT CTTCAGAAAA ATGTGAATGG 60
15 CTTATAAACA TAAATGTAAT ACTCAGCTGC TGTATCATC AGAAGAATGA AGTTGTTTT 120
AAGCTCTTCA NCTGTCAAGN GTAAGAAGCA ATGTTACTAA TACCGGCAAG NTATGAAAAG 180
ACGCTCATAC ATTGTTGGAA ATNAANGTTG GTATNGCTTT TCAGCGGCCN ATNAACATAA 240
TCCCNNTNTT TCTCCCTAC NNACTCN 267

20 SEQ ID NO:6791
SEQUENCE LENGTH:189
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS08019
SEQUENCE DESCRIPTION:
GATCATCTGG CAAGACAGCC CAAGAATAAA GCGTCACCA TGGAGACTAA GAAGNGNCAG 60
AAAGNCACTG GGTCTTCATT GTNTCGTCCA GCTGTTCTAG AGGCCTTATC TGAAGCCAC 120
GTGACCTCCG GNCTACTCAC TTACATGCGN GAATAAAAGC TCTATTTTGT TTAGGNCCN 180
30 AAATAANTN 189

SEQ ID NO:6792
SEQUENCE LENGTH:129
35 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08020
SEQUENCE DESCRIPTION:
GATCTGCTGG CAGAGGNCAA TCAAAAACGA CAACAAGCTT CTTCCCAGGG TGAGGGGAAA 60
40 CACTTAAGGA ATAAATATGG NGCTGGGGTT TAACACTAAA AACTAGAAAT AAACATCTCA 120
AACAGTAAA 129

SEQ ID NO:6793
45 SEQUENCE LENGTH:267
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08021
SEQUENCE DESCRIPTION:
50 GATCAGGACC CACCTCCAGT TCTNCTGAAA GTNTGACAGT GTCCAGCCGG TTCTGCAGCA 60

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EP 0 679 716 A1

CTAGGGGAGG GGGCAGATGG TGGTTGCATG GGCTTCCTGG GTCTCCACTC TCCGTCTGGC 120
 CTAAAGGTGA TGTATTTGGT GTTTGGNCCT GCAGTCCCCA CTNTTAAGGC TTAAGGCGGA 180
 TGTGGAACAN CACTTTCTTT CAGGAGTAGG TNGNTNTACT TGTACCTCAT TTAGGGCTTA 240
 5 GAAAGTTTTN CCNCACTN NAAAAAN 267

SEQ ID NO:6794
 SEQUENCE LENGTH:122
 SEQUENCE TYPE:nucleic acid
 10 TOPOLOGY:linear
 CLONE:HUMGS08022
 SEQUENCE DESCRIPTION:
 GATCTACAAA ACAGATAGNG NAAAGTACAA CAGAATAGCT CGGGAATGGN CTCAGAAGTA 60
 15 TCGGATGTAA TTAAGAAAT TATTGGATAA CCTCTACAAA TAAAGATAGG GGNACTCTGA 120
 AA 122

SEQ ID NO:6795
 SEQUENCE LENGTH:56
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08023
 SEQUENCE DESCRIPTION:
 25 GATCCCACTT GTAGATGCTT GCACTCACAC AATAAAAACG TCTGNTATTT GCCAAA 56

SEQ ID NO:6796
 SEQUENCE LENGTH:130
 SEQUENCE TYPE:nucleic acid
 30 TOPOLOGY:linear
 CLONE:HUMGS08024
 SEQUENCE DESCRIPTION:
 GATCCGTGGT TCCCTGCAGA CTGGCAGGAG CAGATTCCAA AGGCACAGGA AGAAGCTTGC 60
 35 AGGTAGAATG TGTCATTAC CTTCTGCGCA TTATACCACA AAAAAGCTGG GNATAAAAT 120
 GCTAACCAAA 130

SEQ ID NO:6797
 SEQUENCE LENGTH:234
 40 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08025
 SEQUENCE DESCRIPTION:
 45 GATCTGCCCC ATGATTGGTT CCTTAGGANC AAGNAATTTA CAAGTAGNAA TTATTCCTTT 60
 CAGAGTACA TGCTGTATTA CTTCAATCCC TATTTTGTG TGTCCATTT TCTTTGGATT 120
 CCCTATTAC ATTGAATCCT NTTTGCCCTT CTGAAACAAT ATTCAGTCAC AAGNGTTTT 180
 GGTATGTTG GTCTTTGTAA CAAATNAAAN TTACCTTATA TCCTTCTGGA CAAA 234

50 SEQ ID NO:6798

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EP 0 679 716 A1

SEQUENCE LENGTH:246

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08026

SEQUENCE DESCRIPTION:

GATCACAAGG TCAGGAGTTC GAGACCAGCC TGGCCAATAT GGTGAACCTC ATCTCTATTG 60
 AAAATACAAA AATCAGCCAG GCACGGTGGC AGGCGCCTGT AGTCCACCTA CTCGGGAGGC 120
 TGAGGCAGGA GAACTGCTTG AACCCAGCGG CAGAGGTTGC AGTGAGCCAA GATTGAGCCA 180
 CTGCACTCCA GCCTGGTGNT AGNGCGNGCT CTGTCTCTAA ATAATNANNT AAANTNCNTT 240
 AANNNN 246

SEQ ID NO:6799

SEQUENCE LENGTH:114

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08027

SEQUENCE DESCRIPTION:

GATCAGTGTT AAAC TAGATT GTATTCATTA CTAGATAAAA TGTATAAAGC TCTCTGTACT 60
 AAGGNGAAAT GACTTTTATA ACATTTTGAG NAAATAATAA AGCATTATC TAAA 114

SEQ ID NO:6800

SEQUENCE LENGTH:274

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08028

SEQUENCE DESCRIPTION:

GATCTTTTGT GCCAAGATGC AGGGAAATNA AGAAAGCTCT TTCCAGCTGT TTTNATGGAT 60
 TCATGGAGTG GGCTCATGTT GGGACCAGCT GACTCTGGGN GGAAGACTGC CTCTCCATCG 120
 CTGTGAGAGA GCCTGAAAGG ACCAACAATG AGNTGTTCTT GGGACCCGCC ATGGGGATGA 180
 TTGCTTCTCT AGGCTNCTGG CTGTTGCTGC ATTCTAAGGN TTAACCTCCT GGTCTNATGG 240
 TAGTGANCTT GAGCTTTTNN ATTCNATAGN AGNN 274

SEQ ID NO:6801

SEQUENCE LENGTH:42

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08029

SEQUENCE DESCRIPTION:

GATCTGTGNA GNAATGAAAT AAAATGGAGA AGAAGACAGA AA 42

SEQ ID NO:6802

SEQUENCE LENGTH:228

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08030

EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCCAGTTT CAAGAATGGG CAGGTAAACG CAGNANGAGG AAAGGAATGT GGAATGAGAA 60
 CTTGGTGGTT CACCGCTGTA CTATTTGTGT AAATGTTTAC GTATGTNATA AGCTACATGT 120
 ATGTAAATNT TGCAATACCC CTAACAGTCG AGTAGTAGTC TCCCTTACAG GAATTTTGA 180
 CGGGGTTTCT CATCATCAAT ACCAAATAAN TATATGTAGG NNTGGAAA 228

SEQ ID NO:6803

SEQUENCE LENGTH:180

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08031

SEQUENCE DESCRIPTION:

GATCATATTC TTGAAATTTT TATAAATATG TATGGAAATT CTTAGGCTTT TTTTACCAGC 60
 TTTGTTTACA GACCCAAATG TAAATATTAA AAATAAATAT TTGCAATTTT CTACAGAATT 120
 GAATACCTGT TAAAGGAAAA TTACAGAATA ANCTTGTGAC TGGTCTTGTT TTACATTAAT 180

SEQ ID NO:6804

SEQUENCE LENGTH:277

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08032

SEQUENCE DESCRIPTION:

GATCATTCCG AGGTCATTG AGGACGCACA CTGGAGAGAA GCCTTATNAG TGTAACAAT 60
 NTGGAAAAAC CTTCACTTGG TNCTCAACGT TTAGAGAACA TGTNAGAATT CACACGCAAG 120
 AGCAGCTCCA TAAATGTGAA CACTGTGGGA AGGCCTTTAC CTCTTCCAGA GCATTCCAAG 180
 GTCATTGAG GATGCACACT GGAGAGAAGC CTTATGAGTG TACACAATGT GGANAAACCT 240
 TCACTNGGGG CCTCAANCCT TACNTAATAA TGTGGTN 277

SEQ ID NO:6805

SEQUENCE LENGTH:273

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08033

SEQUENCE DESCRIPTION:

GATCTACCTC AGTATTCAAG TTCAGTGGGG ACACCAGTGG CTTCAACTT CCTGGTTTCA 60
 TGATATCTTG AGACGCCCTTA CAAATGATGG AGGATTCCAA AGAGTTTTTG TTTATTGGG 120
 TTAATATTTG TTGGTATTTA TGGCATTGGA GATTGAAACT AAGNAATGTT TTAATTTATT 180
 ACCTTTACAA CATTATTTA CATTACATAC ATACATTAC AACATTATT AATTTATNTT 240
 AAAATAGCAT GNATAAGCCA ATTATNGGTT AGN 273

SEQ ID NO:6806

SEQUENCE LENGTH:207

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08034

SEQUENCE DESCRIPTION:

GATCCACTCT TCTCGNCCTC CCAAAGTGTT GGGATTATAG GCGTGAGCAC ANGCGCCCGG 60
 CCTGNNTG ATATTCTTA TAAATGGAAT CATAAGTGC TTNTTCAGT TAGCATAATG 120
 TTTCCAGGT TCATCCAGGT CATAACACAT AATCAGGCTT CATTNTTTT TATTACCANC 180
 TAATATTCTA TGAATATAAA NNNACCN 207

SEQ ID NO:6807

SEQUENCE LENGTH:273

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08036

SEQUENCE DESCRIPTION:

GATCCTGAAG AGATTGAAAA AGAAGAGCAG GCTGCTACTC AGCCTGAGGT TGCAGACTGG 60
 TCTGAAGGTG TACAGGTGCC CTCTGTGCCT ATTCAGCAAT TCCCTACTGA AGACTGGAGC 120
 GCTCAGCCTG CCACGGAAGA CTGGTCTGCA GCTCCCACTG CTCAGGCCAC TGAATGGGTA 180
 GGAGCAACCA CTGACTGGTC TTAAGCTGTT CTTGCATAGG CTCTTAAGCA GCATGGNAAA 240
 ATGGTTGCTG GCAAATAAAC ATCAGTTTCT AAA 273

SEQ ID NO:6808

SEQUENCE LENGTH:73

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08037

SEQUENCE DESCRIPTION:

GATCTCCAAC ATCTTTGTGT TGAAGGGCCT ACAGAGTAAA CTGTGGGACT CAATAAACAT 60
 TCGTTACTTG AAA 73

SEQ ID NO:6809

SEQUENCE LENGTH:285

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08038

SEQUENCE DESCRIPTION:

GATCACAGCA GACAGGGTCG CTGCCCTGAT GGCCTTACA TTCCAGTGGG TCTAATGACC 60
 ATATCTTAGG ACACAGATGT GCCCAGGGAG GTGGTGTAC TGCACAGGAA GTATGAGGAC 120
 TTTAGTGTCC TGAGTTCAAA TCCTGATTCA GGAACACACA AAGCTATGTG ACCTTACACC 180
 AGTCACTTAA CTTGTTAGCC ATCCATTATC GCATCTGCAA AATGGGGATT AAGAATAGAA 240
 TCTTGGGGTT AGTGTGNGA TTAGATTAAA TGTATGTAAG GCACN 285

SEQ ID NO:6810

SEQUENCE LENGTH:62

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08039

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

GATCAGTTTC TGGCACATAA ATGCCTCAAT AAAGATTAA TTACTTTGTA TAGTGCTTTA 60
AA 62

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SEQ ID NO:6811
SEQUENCE LENGTH:237
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08040

SEQUENCE DESCRIPTION:

15

GATCATTGGC TATGACAGAG AACCCGTCTC TCAAGCAGCA GCTCTTCTCC TATGCCATTG 60
TTGGCTTTGC CTGTCTGAGG CCATGGGGCT TTTNTGTTTG ATGGTCGCCT TCCTAATCCT 120
NTTCGCCATG TGAGGNTCCA TGGGGGGTCA CCGGANCTGT TGATACTGCA ACTACACACC 180
ATTNTTGGTG CTGGGGTGTG TTAAGATTAA ACCATTNANC ACAACGTTTC TCTNAAA 237

SEQ ID NO:6812

SEQUENCE LENGTH:215

20

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08041

SEQUENCE DESCRIPTION:

25

GATCAGCAAA TTTAAGAGAA GACCTAGCTA TGTGGGCTGG TTGCTTTCTA TTATCATGGT 60
ATTTGACCAT TTTAGTTTAA ATTCCATGTC AGATAAGTGT AAATAGAAGA GTTTAAAAGC 120
ATGANACATT TCAGANGGTA TCAGTTATAT GATATTCTTT AANCANNTAT GACACATGTA 180
AATCCTCNTG ANTGACAATN CATCTNTNTG TGAAA 215

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SEQ ID NO:6813

SEQUENCE LENGTH:181

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS08042

SEQUENCE DESCRIPTION:

40

GATCTAAAAG CCATATCCTG TCTTTGGTGG TGGCCTGGTA GGAATGGCTT GTNCTAGTTC 60
TTCAAAGGTA AGATTGACCA AAGCAAGTTC TCATTGGTGG TCTGTGTTTA CATCTTCCGT 120
GTCAAGNTAT TTTAAAGCAT GNAAATAAAG CCATTCTTGC CCTTTTTCGT GTGTTTGCAA 180
A 181

SEQ ID NO:6814

SEQUENCE LENGTH:162

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08043

SEQUENCE DESCRIPTION:

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GATCTCAGGC TTCTGCAGTT CTCATGACTC CTACTTTTCA TCCTAGTCTA GCATTCTGCA 60
ACATTTATAT AGACTGTTGA AAGGAGAATT TGAAAAATGC ATAATAACTA CTTCCATCCC 120
TGCTTATTTT TAATTTGGGA AAATAAATAC ATTCGAAGGA AA 162

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EP 0 679 716 A1

5 SEQ ID NO:6815
 SEQUENCE LENGTH:431
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08044
 SEQUENCE DESCRIPTION:
 10 GATCTCAGCG GGGAGCCACG TCTCTTGCAC TGTGGTCTCT GCATGGACCC CAGGGCTGTG 60
 GGGACTTGGG GGACAGTAAT CAAGTAATCC CCTTTTCCAG AATGCATTAA CCCACTCCCC 120
 TGACCTCAG CTGGGGCAGG TCCCCAAGTG TGCAAGCTCA GTATTCATGA TGGTGGGGGA 180
 TGGAGTGTCT TCCGAGGTTT TGGGGGAAA AAAAAATTN TAGCATATTT AAGGGAGGCA 240
 ATGAACCCN TCCCCANCT NTTNCCNGCC CAAATNTGTC TCCTAGNATC TTATGTGCTG 300
 15 GGAATAATAG GCCTTNACTG CCNCTNNGNT TTATAGGCCN ANGGGTCCCG GTTCCCGGG 360
 AACTGGAACC CTTCTGAGG GGGAATCCCG GGGCTCAAAT TACCCCNAA AGGAAAGTGG 420
 CCAAGGCGGT N 431

20 SEQ ID NO:6816
 SEQUENCE LENGTH:231
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08045
 SEQUENCE DESCRIPTION:
 25 GATCTGTTTC TAAAGCGTTA CGCATCTTTN CTGAAGTATG CTGGAGAGCC TGTCCTTTTC 60
 ATTGAACCCC CTGAAAGCTT TGAATTTNAT GCACAGCAGC TAAGAAAATT GAGGGAAAAC 120
 TCTTCTTGAA ATAACCAGGC GATACTTTGT TTTGTATATA TTTGTGATTC TGTGTCTACA 180
 TGTATTTTG AAGTATATCT GAGGGAANNG GGAATGGAAA TTTCTTTAA A 231

30 SEQ ID NO:6817
 SEQUENCE LENGTH:147
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS08046
 SEQUENCE DESCRIPTION:
 GATCCTTCCA GCTGCTCTTC CATAACTTCA TCTCTCCCT CAACACCTGC AACCCTAAG 60
 CCATTTTCCC CCTACCGTC TGTAATTTTG TAATTTTGAG AATGCCAGGT AAATGGAATC 120
 40 AGAGCGGATA ATCTTTGGAG CATGAAA 147

45 SEQ ID NO:6818
 SEQUENCE LENGTH:101
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08047
 SEQUENCE DESCRIPTION:
 GATCATTAT ATTAAAGTTT AAAAGGAGGN TGAGACTGAG TTTTGTAGA GTTATAATNA 60
 50 CATTCAAAAT ATACTTTTTT GCATTGNAA AATGTTAANT N 101

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EP 0 679 716 A1

SEQ ID NO:6819
 SEQUENCE LENGTH:236
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08048
 SEQUENCE DESCRIPTION:
 GATCAAGCCA AACAGTAAAA ACTACCAAGA GAACACGAGG AAGGCAGAAA CGNTGTTTAG 60
 10 CAACAGTATT CTGCATGGTT CACTGCTTAA GAAAATGCCT TCTGGAATAT TTGTAAACTG 120
 AAATTCTGTA TGTGTAAGNG NTGTTTAAAT ATGTTTGGGT CCTAAACAGC TTTTAAAAAT 180
 TATACTTGGG CNTAANTTCA GCATCTTTTC AAATCANCTC TNTGCNCCGC CGCTGN 236

SEQ ID NO:6820
 15 SEQUENCE LENGTH:195
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08049
 SEQUENCE DESCRIPTION:
 20 GATCATTTTG TTTGTTGCTC TCTCTAGTGT CTTCTTCTCT CGTCAGTCTT AGCCTGTGCC 60
 CTCCCCTTAC CCAGGCTTAG GNTTAATTAC CTGAAAGATT CCAGGAAACT GTAGCTTCCT 120
 AGCTAGTGTG ATTTAACCTT AAATGCAATC AGGAAAGTAG CAAACAGANG TCAATAAATA 180
 TTTTAAATG TNAAG 195

25 SEQ ID NO:6821
 SEQUENCE LENGTH:160
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 30 CLONE:HUMGS08050
 SEQUENCE DESCRIPTION:
 GATCATAAAT ATTTTGTAAT GAAATACTTC CNCTTTTCCA GGGCTTTGTA TGCACTTGTA 60
 TAATTACATT GATGGCAATG TAGNGTTTGA ATTTCACTCT GTAAATACTT TTNTGGNAAA 120
 35 TAGNAATTTT NATTGCTTTA AAGTTTGGGA TATGGGTGGN 160

SEQ ID NO:6822
 SEQUENCE LENGTH:157
 SEQUENCE TYPE:nucleic acid
 40 TOPOLOGY:linear
 CLONE:HUMGS08051
 SEQUENCE DESCRIPTION:
 GATCTATAGG AAGATTGAAC CTGAATATTG CCATTATGCT TGACATGGTT TGCAAAAAAT 60
 45 GGTACTCCAC ATACTTCAGT GAGGGTAAGT ATTTTCCTGT TGCAAGAAT AGCATTGTAA 120
 AAGCATTTTG TAATAATAAA GNATAGCTTT AATGAAA 157

SEQ ID NO:6823
 SEQUENCE LENGTH:150

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EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08052
 SEQUENCE DESCRIPTION:
 GATCCAGNTT GTTGATCAA TAGTCTGTT TTTAGTACTG AGTGGGATGT CAGTATAGAT 60
 GTACCATAGT AAACACCTGT GGAAGGACGT CAGGAATGTT TCCTGTTATT AGGTATCTAA 120
 ATAAAGTTGC TGTAACATT TGTGTACAAA 150

SEQ ID NO:6824
 SEQUENCE LENGTH:268
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08053
 SEQUENCE DESCRIPTION:
 GATCTGTCCA CTTTGCTCTC ACGGACGTGG CATTGCCAG GAAGGAGAAC CCAGCAGGAA 60
 CGGAGGTCTC TGCCTTCACC TACAGTGAAG GGGCTGGTAT GTGAAATACT TTCCTGAGC 120
 TTAGTGAAAA ATTCATTTA CACCCTAGTT AGCTCCAGA AAGGTTNAAC TTTCAAGANG 180
 GTCAAAATAT GNAGCCTGTT ATATGTGTAA TTAAGGAGN NNTAGNATAT CCCAAATGGN 240
 ACATTTTCAGG NATTAANTAA TTGNAATN 268

SEQ ID NO:6825
 SEQUENCE LENGTH:197
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08054
 SEQUENCE DESCRIPTION:
 GATCTGAGAA GTGAGTGTA GGAGTTTAAA TCAACCTCTG TTCTGTGCTA GTTAAGGTAA 60
 TAAAGTTGCT GCAGTCCAGG GGGTAGGTAC CTGTGGACGG CTCTGCGAAT AGGNCAGTTG 120
 CATTTCTTGG GAATCAAGTG CATCCCTAGG CTGGCAGTGC AGCAGAAATA CTGAATAAAA 180
 TGTGACAAAT CTCCAAA 197

SEQ ID NO:6826
 SEQUENCE LENGTH:196
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08055
 SEQUENCE DESCRIPTION:
 GATCAAGTAA AAATGAAGGA CAACGTGAAG TTAAAGTTC GATGCAGCAG ATACCTTTAC 60
 ACCCTGGTCA TCACTGACAA AGAGAAGGCA GAGAACTGA AGCAGTCCCT GCGGCGGCT 120
 TTGGCAGTGA AGGAACTGAA ATNAACCAGA CACACTGATT GGAAGTGTAT TATATTAAAN 180
 TNCTAAAAAT CCTAAA 196

SEQ ID NO:6827
 SEQUENCE LENGTH:213
 SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08056

SEQUENCE DESCRIPTION:

5 GATCATCAGT ACTGTTGTCT CATGTAATGC TAAAACTGAA ATGGTCCGTG TTTGCATTGT 60
TAAAAATNAT GTGTGAAATA GAATGAGTGC TATGGTGTG AAAACTGCAG TGCCGTTAT 120
GAGTGCCAAA AATCTGTCTT GAAGGCAGCT ACACTTTGAA GTGGTCTTTG AATACTTTTA 180
ATAAATTTAT TTTGATAAAT AATATTGAAC AAA 213

10

SEQ ID NO:6828

SEQUENCE LENGTH:133

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

15

CLONE:HUMGS08057

SEQUENCE DESCRIPTION:

GATCGTCCTG CAGGTATTTT GCAAGCCCCT ACTGTCCGCC GACATCAGCG TGAATCTCAC 60
ATGGTGCACT TGGCTCCCTG GGGGAGTNTT NGCNTCTCCT GTTGTCTTTT CCAAAAACCA 120
NTAAAAACCA TGN 133

20

SEQ ID NO:6829

SEQUENCE LENGTH:269

SEQUENCE TYPE:nucleic acid

25

TOPOLOGY:linear

CLONE:HUMGS08058

SEQUENCE DESCRIPTION:

30 GATCTGCTAC ATTTTAAAG TGGGCATGAA AATCTTCTGT TATGTATGTA CTGTCATTAA 60
TTCTGTTATT TTTNCCACT AGTACTATCA ACTACAGTAA AGACCTTCCT CTGGCCCAAG 120
GAATCAAGTT TCAGTAAAGG TGTTTATAGAT GANCATCCCT TAATTTGAGG TGTCCAGCA 180
GCTGTTTTTG GNGANGACAA AGNAAATTAA NGTTTCCCT GAATAANTGC ATTATTNTGA 240
CTGTGACAGT GNCTAATCCC CCTATGGTN 269

35

SEQ ID NO:6830

SEQUENCE LENGTH:297

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

40

CLONE:HUMGS08059

SEQUENCE DESCRIPTION:

GATCTATCCA ATAAAGGAGA GAAACAACCG CACACGCCTG GCTCTCATCA TATGCAATAC 60
AGAGTTTGAC CATCTGCCTC CGAGGAATGG AGCTGACTTT NACATCACAG GGATGAAGGA 120
GCTACTTGAG GGTCTGGNCT ATAGTGTAGA TGTAAGAG NATCTGACAG CCAGGGATAT 180
GGNGTCAGCG CTGAGGNCAT TTGNTACCAG ACCNGNGCAC AAGTCCTCTG NCAGCACATT 240
CTTTGGTACT CATGNTCNT GGCATCCTGG NGGGGTNTCT TCGGGAATT GTTCTGN 297

45

SEQ ID NO:6831

SEQUENCE LENGTH:174

50

SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear
 CLONE:HUMGS08060
 SEQUENCE DESCRIPTION:
 5 GATCATGGTG GTTTGGCAGC AGGGAATTG TCTTGTTGGA GCCTGCTCTG TGCTCCCCAC 60
 TCCATTTC TC GTCCCTCTG CCTGGGCTAT GGAAGTGGG GATGCAGATG GCCAAGCTCC 120
 CACCCTGGGT ATTCAAAAC GGCAGACACA ACATGTNGCT CCACGCGGCT CAAA 174

10 SEQ ID NO:6832
 SEQUENCE LENGTH:253
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08061
 SEQUENCE DESCRIPTION:
 15 GATCAATTTT GGAGACCTTG GCCGCCCAGG ACGTGGCGGC AGGGGAGGAC GAGGTGGACG 60
 TGGGCGTGGT GGGCGCCCAA ACCGTGGCAG CAGGACCGAC AAGTCAAGTG CTTCTGCTCC 120
 TNATGTGGAT GACCCAGAGG CATTCCCAGC TCTGGCTTAA CTGGATGCCA TAAGACAACC 180
 20 CTGTTTCCTT TGTGAACCTT TCTGTTCAAA GCTTTTGCAT GCTTAAGGAT TCCAAACGNC 240
 TACTGTAATT AAA 253

25 SEQ ID NO:6833
 SEQUENCE LENGTH:86
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08062
 SEQUENCE DESCRIPTION:
 30 GATCATGAAC CAAAGGAATT TATGTTTTGT AACTTGGGTA CTTATTTTG CATTTTGTTA 60
 TACTATTAAA TANTTTTNTC CTGAAA 86

35 SEQ ID NO:6834
 SEQUENCE LENGTH:124
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08063
 SEQUENCE DESCRIPTION:
 40 GATCCAAGTT TCACTTTTGT TGATATAATG AGCCAGTTTT CTCAGTAATG TCTACTAAAA 60
 AAAAAGCCAT TAACCNGGNT CTAATATTG CATGATTCTA ATAAANGTAT GTCAACTTCT 120
 TAAA 124

45 SEQ ID NO:6835
 SEQUENCE LENGTH:107
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08064
 SEQUENCE DESCRIPTION:
 50 GATCCCTCA GAAGGCAGGA GTGCTGCCCT CTCCATGGT GCCCCTGCCT CTGTGCTGTG 60

55

EP 0 679 716 A1

TATGTGAACC ACCCACGNGA GGGNATAAAC CTGGCACNNG GACCAAA 107

SEQ ID NO:6836

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08065

SEQUENCE DESCRIPTION:

GATCCTCCTG TCTTGGTTTC CCAAAGTGCT GGGATTGCAG GTGTGAGCTA CTGCACCCGG 60
CCCAATAAAA AATCTCGTT CTGTGAAAGA AATGTTTAAT AAAAGTTGTA TTGTCAAATT 120
TTAAA 125

SEQ ID NO:6837

SEQUENCE LENGTH:178

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08066

SEQUENCE DESCRIPTION:

GATCACTAAG ATTAGCTATA TCTATACAGT CATTAGTTTG ACAAGAAATA GNATCCTGTC 60
AGATGCCAAA GAGTGGGATT TTATGTTTA ATGATTAAAC ACCATTATTT ATTGACAATT 120
TACCCTGTGG AACTGTATTA TTTCTAACTA TGAAATAAAG GGGTGATGTA AACACAAA 178

SEQ ID NO:6838

SEQUENCE LENGTH:222

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08067

SEQUENCE DESCRIPTION:

GATCCCCGGC CAAGCCCAGT CACGNGAATT TAAGGCTCTT CTCTCCGGCT CCAGCCTGTG 60
CCTGTAATTG TCCAGGANTG GCCTAGGCCG ACACCTTCTG CTGCCCCGGA GTCCCCCTCC 120
TGGGTGGGAT GCTCACTACA GCTCTGATGT ATATAGCACA CAGCCCAANT CCGACCCTCC 180
GGTCTGTGCG CAATAAAGNG GGGTGTGTTT CANTCCGGGA AA 222

SEQ ID NO:6839

SEQUENCE LENGTH:198

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08068

SEQUENCE DESCRIPTION:

GATCCCAACA GGCCTGGCA GCGTCTGGAC ATGTGTAAAC AGCAGCAGTC ACGTTCCTCG 60
GCTGNCACAA CCAACACNNN AGCAGGTTTT CCATTTTCTG TACTTTNATA TTTCTGTTCA 120
ACCTGTTGGT TTCTACAATG ATTTTAAACA TTGGNAAGNC AGCCTTGNGT ATATTTTAA 180
AAATTATATT NAAAATGN 198

SEQ ID NO:6840

EP 0 679 716 A1

SEQUENCE LENGTH:284
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08069
 SEQUENCE DESCRIPTION:
 GATCATGAAA CTTAGTAGAG GGGATTGTGT GTATTTNATA CAAATTTAAT ACAATGTCTT 60
 ACATTGATAA AATTCTTAAA GAGCAAAACT GCATTTAATT TCTGCATCCA CATTCCAATC 120
 ATATTAGANC TAAGNTATTT ATCTATGAAG ATATAANTGG TGCAGAGAGA CTTTCATCTG 180
 TGGATTGCGT TGTTCCTTAG GGTTCCTAGC ACTGNTGCCT GCACAAGCAT GTGNATATGT 240
 GANATNNNAT GGATTCTTCT ATAGCTAAAT GNGTCCCTC TNGN 284

SEQ ID NO:6841
 SEQUENCE LENGTH:114
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08070
 SEQUENCE DESCRIPTION:
 GATCCACCTG CTCGGCCTC CCAAAGTGCT GGGATTACAG CTGTAAGTCA CTGATNCCGG 60
 NTGCATGTCA AGCACATTGG NAAGTCTTA CAACAATTCT GNTGGAGGNT TTTN 114

SEQ ID NO:6842
 SEQUENCE LENGTH:253
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08071
 SEQUENCE DESCRIPTION:
 GATCTGTCTC TCAAAGGAAG AAACAAAAAC CAAAATTTT GACTGCAAAA GATTGACTAA 60
 TTCTAGGTGC AAATAACTGG AAATTTAGGA TTGTTATTGT ACTACACAGG TCACAANGTT 120
 CCAAACCTAT TTCCCTTGTA ATTTTATGTG CATGATTTT CAGAATGCAG TTTCTATCTT 180
 TTTAGGTGCA ATATTACGAA CTGCTTAGGT CAGANCANTT TNGTNTCTCC AGTTTTTTCT 240
 CCCTANGTTA CAN 253

SEQ ID NO:6843
 SEQUENCE LENGTH:103
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08073
 SEQUENCE DESCRIPTION:
 GATCTTATAA CCTCATTGTT TCCTAATCAT TTTATTTTCT CTCCTGCTAG TACTGTGCCG 60
 CTTCCCCCTC CCCCCACACA AAATAAAAAC AGTATCTGCC AAA 103

SEQ ID NO:6844
 SEQUENCE LENGTH:110
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS08074

SEQUENCE DESCRIPTION:

5 GATCCGTGCC CCACCTCTCC CCAGTTAAGT GCCTTCACAC AGCACTGGTT TAATGTTTAT 60
AAACAAAATA GAGAAACTTT CTTATAAAT AAAAGTAGTT TGCACAGAAA 110

SEQ ID NO:6845

SEQUENCE LENGTH:246

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08075

SEQUENCE DESCRIPTION:

15 GATCAGGGCT GAGGGTAAGG AAAAGAAGAG ACTAGGAAAG CTGGGCCCAA AACTGGAGAC 60
TGTTTGCTT TCCTGGAGAT GCAGAACTNG GCCCGTGGAG CAGCAGTGTC AGCATCAGGG 120
CGGAAGCTTT AAAAGCAGCA GCGGCNTGTG CCCAGGGCA CCCAGAATGA TTTCCCTATG 180
GGCACCAGGC CAGNGAAAAA ATTGGGCAGC NTCTTTAAAG GGNGGAAAAAT TGTTTGTAGC 240
CCCAAA 246

20

SEQ ID NO:6846

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

25

CLONE:HUMGS08076

SEQUENCE DESCRIPTION:

GATCTACCTG CATATAACAG TCTCTAAGTT TTTTITTCAT TCTTGCANTT TGATAATTTT 60
GATTTTGGCC ANATTAGNNT AATTGGGAC TATCCTGAAA CAGNN 105

30

SEQ ID NO:6847

SEQUENCE LENGTH:26

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35

CLONE:HUMGS08077

SEQUENCE DESCRIPTION:

GATCAAAAGA GCTGTCTGAC TGCAAA 26

40

SEQ ID NO:6848

SEQUENCE LENGTH:84

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

45

CLONE:HUMGS08078

SEQUENCE DESCRIPTION:

GATCCCTTGA GCCCAGGAGT TCCAGACCAG CCTGGGAAAT ATAGGGAAAC CCTGTCTTTA 60
CAAAAAAAAA TTTAAAAAT TAAA 84

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SEQ ID NO:6849

SEQUENCE LENGTH:217

55

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 5 CLONE:HUMGS08079
 SEQUENCE DESCRIPTION:
 GATCAGATTC CTGACTAACC CCTCTCTTAG AGCTACAGCG AGCTGCATTA CCAGCTTAAA 60
 ACACTTCTTA GGGATTAAAT ATAGNTGTNA TTTTCAAAA TCGTTTTTAA TTAAACTGT 120
 GTTTTAGTGT AAAATTGTTA ACCTTGTAAG ATGGATANTG TGTATANGAN TGTAGGCCTT 180
 10 AACTATTTC A CATGAGTCAA ANCANGGCNG CTTTAAAN 217

SEQ ID NO:6850
 SEQUENCE LENGTH:93
 15 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08080
 SEQUENCE DESCRIPTION:
 GATCTCACCC AGTGTGCAAT TTGTGAAAAT GAACCAAATG TCATTACTTT TGTCTTAAAG 60
 20 CAAAGTCTAT TTCAAGNATA TGNACAGCAN AAA 93

SEQ ID NO:6851
 SEQUENCE LENGTH:73
 25 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08081
 SEQUENCE DESCRIPTION:
 GATCAATGAA CCATCTCTTC TGGGCAGTTT TGTTGAAAAT AAAGGTTTCT CTTTGATTTC 60
 30 AAGAATGACC AAA 73

SEQ ID NO:6852
 SEQUENCE LENGTH:104
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08082
 SEQUENCE DESCRIPTION:
 GATCACCGGT TTCCGACTGA GTCTGGGGAT TTNGGCCTTG TTGAACCTC CTACGGTGCC 60
 40 CNGGGAATTT GAAANACGCT TTCTGCATNG ATATNTGGGC CCTN 104

SEQ ID NO:6853
 SEQUENCE LENGTH:123
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08083
 SEQUENCE DESCRIPTION:
 GATCAACATC ATTAAAGATT ATGTAAAAAG TTTAAAGGCT TATGAGCCTA AGTTTGTTCC 60
 50 TATATTACCA TATTTACTGA ATNTTCTGGN AAAGNAACTT TANTAAAGTT TAATCTNAGG 120
 AAA 123

55

EP 0 679 716 A1

5 SEQ ID NO:6854
 SEQUENCE LENGTH:91
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08084
 SEQUENCE DESCRIPTION:
 GATCACTTGA GCCTAGGAGG CAGNGGTCA AGTGAGCTGA GATGGCACTC CTGCGCTCCA 60
 GCCTNGGTGA CAGCGTGAGA NNCTGTCTCA N 91

15 SEQ ID NO:6855
 SEQUENCE LENGTH:219
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08085
 SEQUENCE DESCRIPTION:
 GATCTGTCAG NACCAATTGA AAACCTTGCTC AAGAGCAGTT GCTTATATAT AGTAGGATTT 60
 TACTTTTTC TGCTTATGTA NTACTATATG CTAAAAAAC CTGGAGGAAT ACTTACCAAA 120
 GAGGAGTAAC CATCTCTGAG GGTGGGATTC TGGGGGANTT TTTGTTTTT TCTGTTTTCT 180
 ATANTGTGCA NCTNTTNTAG TATGNTNTT NCTAATTGN 219

25 SEQ ID NO:6856
 SEQUENCE LENGTH:66
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08086
 SEQUENCE DESCRIPTION:
 GATCTTNTC CATTTACAGT ACATTCTTAG GATGTATGTA GTAAATAAAG CTTTCTTTAA 60
 AGCAAA 66

35 SEQ ID NO:6857
 SEQUENCE LENGTH:43
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08087
 SEQUENCE DESCRIPTION:
 GATCACCCCG TAGGAAATAA AGCAGGCATC TCTGGACCTC AAA 43

45 SEQ ID NO:6858
 SEQUENCE LENGTH:338
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08088
 SEQUENCE DESCRIPTION:
 GATCAAAAGA CAGNTAAATN TTTCTTCTAG AACACAGTTA CCCCCTTGCT TCATCTATTG 60

55

EP 0 679 716 A1

5 CTAGAACTAT CTCATTGCTA TCTGTNATAG ACTAGTGATA CAAACTTTAA GAAAACAGGN 120
 TAAAAAGATA CCCATTGCCT GTGTCTACTG ATAAAATTAT CCCAAAGGTA GGTGGTGTG 180
 NTAGTTTCCG NGTAAGACCT TAAGGCCACA GCCAANCTCT TAAGTACTGT GTGNCCACTC 240
 TTGTTGTTAT CACATAGTCA TNCTTGGTTG TACTATGTGA TGGTTAACCT GTAGCTNATN 300
 ANTNIACCTA TTATTCTNTT CCTCCTTTAC CTCANCNN 338

10 SEQ ID NO:6859
 SEQUENCE LENGTH:251
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08089
 SEQUENCE DESCRIPTION:
 15 GATCGTTTGC AAGTAACTGA ATCCATTGCG NCATTGTGAA GGCTTAAATG AGTTTAGATG 60
 GGAAATAGCG TTGTTATCGG NTTGGGTTTA AATTATTGA TGAGTTCCAC TTGTATCATG 120
 GCCTACCCGA GGAGANGNGG NGTTTGTAA CTGGGCCTAT GTAGTAGCCT CATTTNCCAT 180
 CGTTTGTATT ACTGACCACA TATGCTTGTC ACTGGGCANG AAGCCTNTT CNGCTGCCTN 240
 20 CACGCANTTT N 251

25 SEQ ID NO:6860
 SEQUENCE LENGTH:359
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08090
 SEQUENCE DESCRIPTION:
 30 GATCTTAATA TAAATNCACT TTCATTTTTN ATAGCTGTCC CATCTGGTCA TTTGGTTGGC 60
 ACTAGACTGG TGGCAGGGGC TTCTAGCTGA CTCGCACAGG GNTTCTCACA ATAGCCGATA 120
 TCAGAATTTG TGTGAAGGA ACTTGTCTCT TCATCTAATA TGNTAGCGGA NAAAGGAGAG 180
 GAAACTACTG CCTTTAGAAA NTATAAGTAA AGTGATTAAG GTGCTCACGT TACCTTGACA 240
 CATAGTTTTT CAGTCTATGG GTTTAGTTAC TTTAGNNGGG NAGCNTGTNA CTTATATTAN 300
 TAGTANTTTN GTAAAGTTGG NTGGATAAGC TATCCATGTT GCCAGGTTCA TGGATTNGN 359

35 SEQ ID NO:6861
 SEQUENCE LENGTH:75
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 40 CLONE:HUMGS08091
 SEQUENCE DESCRIPTION:
 GATCCCCAAA CCACAAATAC ATATGTGTGC CTAAGTTTGT GCATGCATAC ATTAACAGG 60
 ACATAAAGTG TAAA 75

45 SEQ ID NO:6862
 SEQUENCE LENGTH:228
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 50 CLONE:HUMGS08092

55

SEQUENCE DESCRIPTION:

GATCATCATG GCAAAACCAG CTGGTGGGCC CAAGCCTCCA AGTGGGAAGA AAGACTGGGA 60
TGATGACCAA AATGATTGAA ATTGGCTTAA TTTTACTGT AGGTGAAGGC TGTATTGTA 120
GTAGTACTCA AGAATCACCT GATGTNTTCT TATTCTCCTT AAATTAAGAG TTATTTTGTN 180
TTTGTNTTCT TGGCTGGATG TTATAATAAN CATATTGTNA CTGTCAAA 228

SEQ ID NO:6863

SEQUENCE LENGTH:299

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08093

SEQUENCE DESCRIPTION:

GATCATGCCT ACTTCTTCT NTGTATCTCC CTCCTTCCCA GCCCACC CGG GCAGCAGACT 60
CTNATGGAAG NAAGGTGCCG TAGTGGGNTT TTAGAACTA ACGGGACTGG TTTTCAAAGC 120
AGTTATCTTG GGAACTGTT TATTCCAGCG ATGTGACTTT TTNAGAATA TTTCTTGGAA 180
TCATATTGAG AGTCTGGGGC TGTGTGTTGA GCAGCCTTAA GGNTGCTAGA CACTCATTTA 240
GTGCCCAGGG AGTCCAGCGA ATGACGCTCG TGGCCAAGCG AGGTCTCAGG TGCAAAGCN 299

SEQ ID NO:6864

SEQUENCE LENGTH:298

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08094

SEQUENCE DESCRIPTION:

GATCCTCTGA ACCTGCAGAG GCCCNCTCCC CGAGCCTGGC CTGGCTCTGG CCCGGTCCTA 60
AGCTGGACTC CTCCTACACA ATTNATTNA CGTTTTATTT TGGTTTTCCC CACCCCTCA 120
NGNNTGTGCG GGAGCCCCTG CCCTTNACCT AGCTCCCTTG GCCAGGAGCG AGCGAAGTGT 180
NGGCCTTGGT GAAGCTGCCC TCCTCTTCTC CCCTCACACT ACAGCCCTGG TGGGGGAGAA 240
GGGGGTGGGT GCTGCTTG TGTTAGTCTT TTTTTTTTTT TTTTNTTN TNANAAAA 298

SEQ ID NO:6865

SEQUENCE LENGTH:296

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08095

SEQUENCE DESCRIPTION:

GATCTGCGGG GCAANTTCAA GCNNNCCCCA CTGCGGAGGG TGCGCATGTC GNCCGATGCC 60
ATGCTCAAGG CCCTGCTGGG CTCGAAGCAC AAGGTGTGCA TGGACCTGAG GGCCAACCTG 120
AAGCAGGTCA AGAAGGAGGA CACAGAGAAG GAGCGGGACC TGCAGACGT GGGTGACTGG 180
AGGAAGAACA TCGAGGAGAA GTCTGGCATG GAGGGCCGGA AGAAGATGTT TGANTCCGAG 240
TCCTAGGCCA CTCGCTGCCC CTACGNCTGC CCCGGTGCCC GGTTCACAGC AGAANN 296

SEQ ID NO:6866

SEQUENCE LENGTH:310

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08096

SEQUENCE DESCRIPTION:

5 GATCCTGTTT TGTCAGCTGC ACCCTGAGCA AAAACTTGCC ACACCAGNGC TCCTAGAAAC 60
 AGCCCAAGGCT TTGGAGCGGA CCCTGAGGAA GCCTGGGAGG GGTGGGATG ACACGTATGT 120
 CTTGTCAGAT ATGCTCAAGA CGGTGAAAAT TGTGTGACTG CTTAGGCCAA GCAGCCCTCC 180
 TGCCTAGAAT GACCTTGGAC TCCCAGCCTG CCAGNGAATG NAGAAATACA ACGCACAGTA 240
 10 CTTTGAAGC TTCGTATTTT TCTTGGTTTC AACTCAGCT ACATGTGACC TCCAGCTTGG 300
 TCGGTTAGN 310

SEQ ID NO:6867

SEQUENCE LENGTH:285

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08097

SEQUENCE DESCRIPTION:

20 GATCAGTTCA AGAAGGTAAT CCAAGGGCCT GATGACTCTT TTGGTAACCA GACACCAATC 60
 AAATAAGGGG AGGAGATGAA AATGGAATNA TTTCTTCCAT GCCACCTGTG CCTTTAGGAA 120
 CTGCCAGAA GAAAATCCAA GGCTTTAGCC AGGAGCGGAA ACTGACTACC ATGTAATTAT 180
 CAAAGTAAAA TTGGGCATTG CATGCTATTT TTAATACCTG GNTTGCTGAT TTNCAAGAC 240
 AAAATACTTG GGGTTTCCA ATAAAGATTG TTGTAATATT GNAAG 285

25 SEQ ID NO:6868

SEQUENCE LENGTH:201

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

30 CLONE:HUMGS08098

SEQUENCE DESCRIPTION:

35 GATCAGACTC TGTATCAAAA GTACCTTTGC CCTTAGGAAG AGTGAGTATT GGAGTCATCT 60
 TATCTATTAC TCCAAACCTC CCTTTNATT TCTTGAGCCT GGCTTGGACC TTGGCATTCC 120
 GTTTGANTTC CTTCTAACTG GAACATTTGT GTTGATCTG TAACACTGGC ACTGAAATAA 180
 AGACCACACG GTTAAAGNAA A 201

SEQ ID NO:6869

SEQUENCE LENGTH:50

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08099

SEQUENCE DESCRIPTION:

45 GATCATGTGC TGCCTTTTAC TGCTGAATTA AAACAGATAT TTCACGNAAA 50

SEQ ID NO:6870

SEQUENCE LENGTH:43

50 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08100

SEQUENCE DESCRIPTION:

GATCTTCAGT GTAAATNACC ATGTATAAAC TGTAAAGTGC AAA

43

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SEQ ID NO:6871

SEQUENCE LENGTH:247

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

10

CLONE:HUMGS08101

SEQUENCE DESCRIPTION:

GATCATTAA TATATTCATA TTACCAAGAC TATTATACTG GAAGTGGTTT TTGTGTTATA 60

AAGGTTTAAT TTTACATANG GCAGTTACTT AATGTGATTT TNAACCCCTA AAAAAGTGGA 120

15

GATGTATACA TTTGTTAACA ATGCCATGAA AGCATTNCT TTCTCCTAAG GAAAAGTGAA 180

TTTCTTATCA GAATATCTGG CTGGCCCTGT AATTTAAATT AAAATAAANT TTTGGCGNAN 240

CAGCAAA 247

SEQ ID NO:6872

20

SEQUENCE LENGTH:288

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08102

25

SEQUENCE DESCRIPTION:

GATCTATCTG AGAACAAAGA AGTCGACAGA ATAACATTGA TGACTTTTTA AATAATTAAT 60

TTATTTGAAG AATGCTTGCA TACAAAAAGC TTCCTGCAGG AATTTTAAAC AAGCTGTCCT 120

TGGTTATCCC AGGCAATGTC AGTTGATTTA NTAAAGTTTA CATGCAAAAG NGAATACAAT 180

TGTAACATTA ATTGGAACAT TAAGTTGGCT TATAAGCAGT ATTTCCACTG CTCCTACAAT 240

30

TGNTTTNNGT TATTTGATAN TTNATTTNAN ATTNNGTNTT GGTGTGGN 288

SEQ ID NO:6873

SEQUENCE LENGTH:294

35

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08103

SEQUENCE DESCRIPTION:

GATCGTACCT ATTGCACTCC ATCCTGGATG AAAGAGCCAG ACTCTGTCTC AAAACAAACA 60

40

AAAAAGCGTG GGGACTTCTG GGGACAGACA AGGTGCCTGT AATATATTTN CTCAGTCTTT 120

GCCCTGAATG GTCTCAGCTT GANACCATNT CAAACTGGAG AGAAGCAAGC CAGCCAATAG 180

AATGGGGTGA TTTACAGGGA TTTCTGTTTA CTGTCAAAAT ATTTCTCATC TGCATATGT 240

TTCCATTTGT GGTCTGAAG GAAATTCTTA TAACTCAACA TTTGTCTGGT CTTN 294

45

SEQ ID NO:6874

SEQUENCE LENGTH:295

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

50

CLONE:HUMGS08104

55

EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCCGACAC TCCAGGCAGA GCAGAGGGCA GGAGAGGCC AAAGAGCTAG GTCAAGCAGC 60
 TGGCTCCCT GGAAGGGGAG CCGGGAGACA GGGAGGCCT CACTGCCC CCGTGGCCAC 120
 GCATGGCTGT CTGCACACTC CATCAGGAGG AAGGGCAGGG GAGAAGTCAC ATGCNGTACA 180
 GTGAAACTNA TCAGTGTAC ACACGCCAG CAGGAGGGGA GAGCTCTGCT CCACACGCTG 240
 GCGGCCTTGG CTGCTCCANA GCAATCCGNT TGCTCCANCT TCAAGCCATT GGGGN 295

SEQ ID NO:6875

SEQUENCE LENGTH:168

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08105

SEQUENCE DESCRIPTION:

GATCCTGTAT TATCCATCTA CATCAGAACC AACTACTTC TCCAACACC GGCAGCACTT 60
 GGCCTGCAA GCTTAGGATG AGAACCACTT AGTGCCCAT TCTACTCTC TCATTCCCTC 120
 TTATCCATCT GCAGGTGAAT CTTCAATAAA ATGCTTTTGT CATTCAA 168

SEQ ID NO:6876

SEQUENCE LENGTH:141

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08107

SEQUENCE DESCRIPTION:

GATCCTTGAG ATGACTGTAG CCCTGGCTGA CAACTTNATT GCTGCCTGTN AAGGACCCTG 60
 AACCAGAGGA CTCAGCGAAA TTGTGCCAG ATTCCTGACC TACAGCGCA GTGAGATAAT 120
 AAATGCTGTT GAGGTCACAA A 141

SEQ ID NO:6877

SEQUENCE LENGTH:124

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08108

SEQUENCE DESCRIPTION:

GATCTGCTCA CCCTCGGTNC ACANCAGTGT CAGCCATGCA AGCAGGACAG AATGGTGACT 60
 GGGTGCCCAT NGTGAGCTGT GTATTCCTA GGAGGTAGAA AACAATAAAA AATTGCATGC 120
 TAAA 124

SEQ ID NO:6878

SEQUENCE LENGTH:280

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08109

SEQUENCE DESCRIPTION:

GATCCTCTG CCTCAGCTC CTGGGTGGCT GGGACTGCAG ATACATGCCA CCATGCCGGA 60
 CAAATTTNTT GTATTTNAG TAGAGATGGG GTTTTACCAT ATTGCTCGGA TTGGTCATGA 120

EP 0 679 716 A1

5 ACTCCTGGAT TCAAGCAATC TGCCACCTA GCCTCCCTAA GTGCCTGGCC AAATTTTGG 180
TTTTAATAGT CGATTGTGT ATATTTTGTG GTAGTATCTA CATATATTTA ATACATTTAT 240
NTGTTGNTGN AAAAAGTANA ATTTGTAAAA NNTTTGNAAA 280

10 SEQ ID NO:6879
SEQUENCE LENGTH:198
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08110
SEQUENCE DESCRIPTION:
15 GATCATTGTT GAACTTCAGC TCCAGTGTCT CTCCAGAATA AGACATTGGC ATTCAAATGT 60
CTATATCTTG TTAATTACAA AATAAAAAAC AGATTAATTA GTGGCTTTTA AATTGTAGTT 120
ATATCNGTGT ATATACACGC GGGGNACTGT ATAAAGACAT ACTAAAGGGC ACAGATTAAA 180
ATAAGTATTA TTANTAAA 198

20 SEQ ID NO:6880
SEQUENCE LENGTH:66
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08111
SEQUENCE DESCRIPTION:
25 GATCAAAGTT GTTAAATTA TAAATNTATG ATGCAGNAAT AAAATTGATA TATTTTGATA 60
TTCAAA 66

30 SEQ ID NO:6881
SEQUENCE LENGTH:100
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08112
SEQUENCE DESCRIPTION:
35 GATCTCTTAC CTTTGGAAAA TAGGGGTTAG GCATGAAGGT GGTGTGATT AAGAAGATGG 60
TTTTGTTATT AAATAGCATT AANCTGGAAT TGACAAGAAA 100

40 SEQ ID NO:6882
SEQUENCE LENGTH:30
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08113
SEQUENCE DESCRIPTION:
45 GATCAAAGAA ATAATTCTTT TGCCACTAAA 30

50 SEQ ID NO:6883
SEQUENCE LENGTH:172
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08114

SEQUENCE DESCRIPTION:

5 GATCACTACA ACAATTTGNC TGCCTCCAAG GTCCTCTGAG GCAGAAGGCT CTGGGGNTTC 60
TGCTGTCCCTT TGGNGGGTGT CTTCTGGGTA GAGGGATGGG AAGGAAGGGA CCCTCACCCC 120
NGGCNCTTCT TCTGACCCTG CCACTAAAAA ATTTATGGNC CAAGGGCNCA AA 172

SEQ ID NO:6884

SEQUENCE LENGTH:291

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08115

SEQUENCE DESCRIPTION:

15 GATCAGCAAT NAGGGTGAGG AAAAAATCTT CTTAATCAAC AAGCTTCACT CCATCTACGA 60
GAGGAAGGAG AGGGAGGAGA GGAGCAGGGT TGGGACAACC GAGGAGGCTG CGGCACCCCC 120
TGCCCTGCTC ACAGATGAAC AGGATGCCTA GGGGGACGGC GATGGGCCTC ACGGGCCCGC 180
CCAGCACCCCT GAGACCACAC TGTTCCTCC CAGTGACCCT GCTGGGACAC CAGGACAAGG 240
AAGACAGTTT CGCCTCTNGA AAGCCGCAGC TGGCCTAGG CTGGAGCTGG N 291

SEQ ID NO:6885

SEQUENCE LENGTH:30

SEQUENCE TYPE:nucleic acid

25 TOPOLOGY:linear

CLONE:HUMGS08116

SEQUENCE DESCRIPTION:

GATCTGTAAA TAAAACTTAC ATTTTTCAAA 30

SEQ ID NO:6886

SEQUENCE LENGTH:291

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08117

SEQUENCE DESCRIPTION:

35 GATCTGCCCTG CCTCGGNCTC CCAAGTGCTG GGATTACAGG CGNGAGCAGC TGCACCCAGC 60
CTAGAATCTT GTATAATATG TAATTGTAGG GAACTGCTC TCATAGGAAA GTTTTCTGCT 120
TTTTAAATAC AAAAATACAN AAAAATACAT AAANTCTGAT GATGAATATA AAAAAGTANC 180
40 CAACCTCATT GGNACAAGTA TTAACATTTT GGAATATGTT TTATTAGTTT TGTGATGTAC 240
TGTTTTACAA TTTTNACCAT TTTTTCAGT AATTACTGTA AANNNGNNNN N 291

SEQ ID NO:6887

SEQUENCE LENGTH:288

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08118

SEQUENCE DESCRIPTION:

50 GATCCCAAGC TCAGTCCCA CAAAGTTCAG GGCCGGTCGG AGGCAGGGGC AGGTCCGGGT 60

EP 0 679 716 A1

5 CCAAAGCAAG GACACCACAG CTCTCCGAC TCCAGCAGCA GCTCCAGCGA TTCGGACACG 120
GATGTGAAGT CCCACGCTGC TGGCTCCAAG CAGCAGGAGA GCATCCCGGG CAAGGCCAAG 180
AAGCCCAAAG TGAAGAAGGA GAAGGGCAAG ANGGNGAAGG GCAAGANGAN GGAGGCTCCC 240
CACTGANGGG CCCTGGACAG GGCTCATTA ACCTTCCTCT CTGCCAAA 288

10 SEQ ID NO:6888
SEQUENCE LENGTH:308
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08119
SEQUENCE DESCRIPTION:
15 GATCCGCATT GAAGACCCAC CCCGCAGAAA GCACATGGAA TTCCTGGGTG GTGCAGTTCT 60
AGCGNTTATC ATGAAAGNCA AAGACAACCT TTGGATGACC CGACAAGAGT ACCAAGAAAA 120
GGGTGTCCGT GTGCTAGAGA AACTTGGTGT GACTGTTCGA TAAACTCCAA AGCTTGTTCC 180
CATCATACCC GTAATGCTTT CTTTTTCCT TTATTGCCAA TCTTTGAACT CATTCAACTC 240
CAGGACATGG AAGAGGCCTC TNTCTGCCCT TTGACTGGAA NGGTCAANGT TTTATTCTGG 300
20 TTGTCTGN 308

25 SEQ ID NO:6889
SEQUENCE LENGTH:308
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08120
SEQUENCE DESCRIPTION:
30 GATCATGTCT TTTGCAGGAA CATGAATGGA GCTGGAGGCT ATTATCCTTA GCAAATAAT 60
GCAGGAACAG AAAACCAAAT ACTGCACGTN CTCACCTATA AGAGGGAGCT AAATNATGAG 120
AACTGATGAA CACAAAGAAG GAAACAACAC ACAATGGGGT CTACTTGACA GTGGAAGGTG 180
GGAGGAGGGN GAGGAGCAGA AACATAACT ATTGGATACT GGGCTTAGTA TCTGGGTGAT 240
GAAATAATCT GTACAACAAA CCCCCATGNC ACAAGTTCAC CAATGTAACA NACCTTCACA 300
35 TGTACCCN 308

40 SEQ ID NO:6890
SEQUENCE LENGTH:224
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08121
SEQUENCE DESCRIPTION:
45 GATCAATTCC CTTGAATAGG GAAGTAACAT TTGCCTTAAA TTTTTCGAC CTCGTCTTTC 60
TCCATATTGT CCTGCTCCCC TGTGACGA CAGTGCAATTT GCCTTGTNAC CTGTGAGCTG 120
GAGAGAACCC AGATGTTGTT TATTGAATCT ACAACTCTGA AAGAGAAATC AATGAAGCAA 180
GTACAATGTT AACCTAAAT TAATAAAGA GTTAACATCC CAAA 224

50 SEQ ID NO:6891
SEQUENCE LENGTH:28
SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08122
 SEQUENCE DESCRIPTION:
 5 GATCAAATAA AGTATAAAAA TTGCCAAA 28

SEQ ID NO:6892
 SEQUENCE LENGTH:140
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08123
 SEQUENCE DESCRIPTION:
 15 GATCAGGCCT GGCCACTCCT CTTTGGAGGG CGGTTGAGGC CCACCTGCCT TGGGCTTCCC 60
 AATNCTGATT CCNNTACTAC ACCCTTCCCA GTTCTATTCC CCTCACCACA CTGCTCCAA 120
 TAAAACTATT TATATACAAA 140

SEQ ID NO:6893
 20 SEQUENCE LENGTH:163
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08124
 SEQUENCE DESCRIPTION:
 25 GATCGTCATC CCTTCTGGA GTTCCTATCT TCCAAGATGT GACTGTCTGG AGTTCCTTGA 60
 CTAGGAAGAT GGATGAAAC AGCAAGCCTG TGGATNGAGA CTACAGGGGA TATGGGAGGC 120
 AGGGAAGAGG GGTGTTTCT TTAATAAAT CATCATTGTT AAA 163

SEQ ID NO:6894
 30 SEQUENCE LENGTH:253
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08125
 35 SEQUENCE DESCRIPTION:
 GATCACCTGA GGTGAGGAGT TTGAGACCAG CCTGGCCAAC ATGGTGAAAC GCCATCTCTA 60
 CTAAAATTGA AAAATTAGCT GGGCATGGTG GCAGGTGCCT GTAATCCCAG CTATTGGGGA 120
 GGCTGAAGCA GGAGAATCGC TTGAAACAGG AGGCAGAGAT TGCAGTGAGC CGAGNGGGGN 180
 40 GCCACTGCAC TCCAGCCTGG GCAACAGGAA CAAAACTCTA TCTCAAAAAA TAAACAAGA 240
 TTTTCTGCG AAA 253

SEQ ID NO:6895
 45 SEQUENCE LENGTH:99
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08126
 SEQUENCE DESCRIPTION:
 50 GATCACTTGC AGCCAGNAA GTTTGAGACC ACCCTGACCA ACACAGTGAA ACCCNCTCTC 60
 TAAAAAGTT TGTTTTTAAA GTCTCCTTGA AACATCAAA 99

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SEQ ID NO:6896

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08127

SEQUENCE DESCRIPTION:

GATCAAGCTA TGCCTGATNC TGCCTTAAGC CTTATNAGGT CGCCACGGAA CCGCAGAAAC 60
 TCCTGGTGGG CAGAGCTGTG TGTGCCATCT TCATCATTGC ATTACCACCA CCAAGCACAG 120
 TGCCTGGCAC AAAAGAGTTG CTCAATAAAT AAATCAGGAT GANTGGNTAA ATACATGGNT 180
 AGGCACCTTG AGCTACAGAT GCGTTTAAAT ACTTTGTGTT TTTCTTAGTC AAACATGTGC 240
 ANTTANGCAT GTGATAAATG CTATGATGAC CACAANANTG CACAAA 286

SEQ ID NO:6897

SEQUENCE LENGTH:137

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08128

SEQUENCE DESCRIPTION:

GATCATGGTG AAATATTTTN GATATATAAA TTCCTTAAGC TATTGTAACC ATGTTTTATT 60
 GCAAAGATGT AAAATATGCC AGATGTGTGT NAGTTGGAAA TCAAAAAAAG NAAATAAAN 120
 TATGCAAAGA NTTCAAA 137

SEQ ID NO:6898

SEQUENCE LENGTH:290

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08129

SEQUENCE DESCRIPTION:

GATCCGCAAC AACTCAGCCA TCCACATCCG AGTCTTCAGG GTCACACCCA AGTAATTGAA 60
 AAGACACTCC TCCACTTATC CGNTCCGTGA TATGGCTCTT CGCATGCTGA GTACTGGACC 120
 TCGGACCAGA GCCATGTAAG AAAAGGCCTG TTCCCTGGAA GCCCAAAGNA CTCTGAATTG 180
 AGGGTGGGGG TAATTGTNTG TTGGTNGGCC CAGTTAGTGG GCTTTNNTAA GTGTGTGTAT 240
 GCGGTCTGTA NCTATTGCCA TATAAATAAA NAATCCTGTT GNACTNGTAN 290

SEQ ID NO:6899

SEQUENCE LENGTH:71

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08130

SEQUENCE DESCRIPTION:

GATCATAAAA CGTTGTAAGT TTTCTAAAAAC ACATAAGCTC TCAATAAACG TTAGCTTATC 60
 ATTGTCACAA A 71

SEQ ID NO:6900

EP 0 679 716 A1

SEQUENCE LENGTH:211
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08131
 SEQUENCE DESCRIPTION:
 GATCAAAACAC TGTCAGAATT GCTGAAATCA ATACACAAAG AGATAAAGTT TAGCTTCTTT 60
 TTACTATTCA ATATTGAACA TAATATTGTT AAATATNGNG ATGAAATGCT GTTGGATNTG 120
 CTNCATTAAA TCTTAATGTA ATATTGTAAG NCTTTTGAGA ATATACTTGA TTTAAATGTG 180
 AAAGNNGGGN TTGTTANCTT ATTGCTATNC N 211

SEQ ID NO:6901
 SEQUENCE LENGTH:81
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08132
 SEQUENCE DESCRIPTION:
 GATCATCATC TNCITGCGGA CTTCTCTGCC TGGTCTTGTG TGTCTGTGTA TTCAAACAAT 60
 AAAANGCTGG TNGAACTTAA A 81

SEQ ID NO:6902
 SEQUENCE LENGTH:175
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08133
 SEQUENCE DESCRIPTION:
 GATCAGACAC TGCCTTTTGT GTGTTTGCTG CCTNTGGATN CTTTTTTAAA AAGACTGTTA 60
 CATATTAAAA NAGTGTACAT ATATAAATAT NACCTCTTTT GCTGTACAGT TGTGATAGAG 120
 ACTGAAGNTT TTATTTTNG TGTGCTTTT ATAAGNAAAA ANTTAATACA CTAAN 175

SEQ ID NO:6903
 SEQUENCE LENGTH:92
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08134
 SEQUENCE DESCRIPTION:
 GATCATCTAA TTCTCTGTGG AATAAATACA CACATATATA TTACAAGGGA TAATTTAGAC 60
 CCCATACAAG TTTATAAAGA GTCATTGTGA AA 92

SEQ ID NO:6904
 SEQUENCE LENGTH:312
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08135
 SEQUENCE DESCRIPTION:
 GATCTGAAAA ATACTGAAAA ATGTGATACT TAGAATACTT TGGCTGCTAA GGAAACTTCC 60

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5 TCTCCATTGC AGAATAGCTG AGCCAAGTNA GTGAGTTTNC AGAAAGCAGG TGGTGAGCTC 120
 NTGCCTGCTG GAGGTGCCA TGGAGGGCCA TTCCTGCCCC GCAACAGCAC CGTCCTGCAG 180
 GGAGCCACTT GGCAGAAGGG TGCAGGGCTG CTGGTGTGAG AGCAAGAGGG CTACAGGGAA 240
 AGGGCCCTTT CTCAGGGGAT GTNGNTTTT TAAAAGGTTT GGGNACACTT GCGGGGTTTG 300
 CTAATAATTGA GN 312

10 SEQ ID NO:6905
 SEQUENCE LENGTH:123
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08136
 15 SEQUENCE DESCRIPTION:
 GATCTCTCTT GCAGCTTGAG AATGTCCCAG TTAAGCCACT TAAAAANTCC TCTNTCCTGC 60
 AGCCCAACAC TTTNCTAAGA GGTACAGAAA NGANAGGAAA TAAACACAC ATCATTATGG 120
 AAA 123

20 SEQ ID NO:6906
 SEQUENCE LENGTH:212
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08137
 SEQUENCE DESCRIPTION:
 GATCTCAACA CAGACCCTNA CCAGCTGATN AATNCAGTNA ACACACTGGA CAGGGATGTC 60
 CTCAACCAGC TACACGTACA GCTCATGGAG CTGAGGAGCT GCAAGGNTTA CAAGCAGTGT 120
 AACCCCGGA CTCGAAACAT GGACCTGGGG ACTTAAAGAT GGAGGAAGCT ATGAGCAATA 180
 30 CAGGCAGTTT CAGCGTCGAA AGTGGCCAGA AN 212

35 SEQ ID NO:6907
 SEQUENCE LENGTH:235
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08138
 SEQUENCE DESCRIPTION:
 GATCAAGCCA AAAGAAATAT TTTNAAATA AGCCCTTTTC AAAAGTTTTT GAATTTATAG 60
 40 AAAGCACCAA TGAATAACAT ATTTCTGTTT CGTTAATGTC AGCTGCCTGA ACATTCAGCA 120
 GTTTATAAAT TGCTTAATTT GTGTNATCTA TTATCCAGTA AACCCATAGT TCCATGATAT 180
 GTCACAGGNA TTGTNAGGTC CTATTTTAAA GGTACAGTCT TGTGGNATGT CATCN 235

45 SEQ ID NO:6908
 SEQUENCE LENGTH:224
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08139
 50 SEQUENCE DESCRIPTION:
 GATCCTTTGA GCCCAGAAGT TGGAGACCAG CCTGGACAAC ATGACAAAAC CCCATCTCTA 60

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EP 0 679 716 A1

5 CAGAAAATTT AAAAATTGGC CAGGCGTGGT CGCGCACGCT GTAGTCCCAG CTATTCAAAA 120
GGCTGAGGTG GGAGGATTAT GTCCCTGGAG CCCAGGAAGT GGNGGTTGCA GNGATTGCAT 180
CACTGCATT NAGCCTGGGT GACAGNGTAC GCCNCTGTAT CAAA 224

10 SEQ ID NO:6909
SEQUENCE LENGTH:215
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08140
SEQUENCE DESCRIPTION:
15 GATCCTGCTG GCCAGAAGCA AGCGGGNCG GCAATAAATC CAAGAAATTG TCCCAACAAC 60
CACCAATTCT TACGGAGGAN TATTATTTAG CCAGCAGGAG TCGNGTTTGG TTTACTGCTT 120
TTACTGTTTT GNGTTCATGN CTNTTATTT TAATGGCGTT AAAAGCACAG GCAAATGTCT 180
TTGTAATGC AGCTTAGTAT GANTTCTNTT ANAGN 215

20 SEQ ID NO:6910
SEQUENCE LENGTH:253
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08141
25 SEQUENCE DESCRIPTION:
GATCACAGAC TCTGAAGNCA GACTGGCAAG ATTCAAATCC CGACTGCCAT TTTAACTGA 60
AAGACCTTAG GTAAGTTACC TAACCTCTCT GTACCTGTGT AAAATGGNGA TAACAGTAGT 120
ATCCACTTCA CAGAGTTATG AGGNTGAAAT ACATTCATAT TTATAAAACG TTTGGACCCA 180
GTACTTGCA CATAGTAAGG CCATGTAAAGT GTTTGTAAAA TAAAACTGNA AATAAAGCCA 240
30 TNTCCNAATN AAA 253

35 SEQ ID NO:6911
SEQUENCE LENGTH:359
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08142
SEQUENCE DESCRIPTION:
40 GATCACACTG CTGCAAGACC TGACCAGTTG AGCTTCCGGC GTGGGGAAGT GCTGCGTGTC 60
ATNACCACAN TGGATGAGGA CTGGCTCCGC TGTGGGCGGG ATGGCATGGA GGGTCTNGTG 120
CCTGTGGGGT ATACCNACCT TGTTCTGTAG CCCTGGNGNC CTTTCTGCG TATGTGTCTC 180
CTTCCTGTCA CCTGGGCATG GAATGGCCAG TGAACACCAT CCCAGAAGCA TTTTCCCTCT 240
GCAAAATGAC GTTTCTTCCC ACGTCTGTTT CTGCTAATAT TAAAAATAA CTTTCTTCT 300
45 TCCCTCCTAT ACCANCTGTA AGGGAAAATN TGCTCTTNTT CCAAATATAT AAAAAGGTN 359

50 SEQ ID NO:6912
SEQUENCE LENGTH:49
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08143

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SEQUENCE DESCRIPTION:

GATCTTTTAC ATTGCAAACA AAATGCTGTA TTGCTTAAAA GGGTTGAAA 49

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SEQ ID NO:6913

SEQUENCE LENGTH:30

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS08144

SEQUENCE DESCRIPTION:

GATCAGTTAA TAAACATGAG TAGCTTGAAA 30

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SEQ ID NO:6914

SEQUENCE LENGTH:128

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08145

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SEQUENCE DESCRIPTION:

GATCTGAAGT GTACAAAGTT CTCATAGCAC CNCTGGGAAG AGAAGGACGT AAACAGAACC 60

TGACACCAGC TCCTTTTCCT TNTATACATT ATTTAATACC TATTAAATAC AATTATTTTT 120

GGAATAAA 128

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SEQ ID NO:6915

SEQUENCE LENGTH:425

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS08146

SEQUENCE DESCRIPTION:

GATCTACTTG TAAGTTGTTG GCCTTCTTCC CACATCTGCC TCANACTGGG GGGGGCTCAG 60

CTCCTCGGGT GATATCTAGC CTGCTTGTTA GCTCTAGCAG GGATAAGGAG AGCTGAGATT 120

GGAGGGAATT GTGTTGCTCC TGGAGGGAGC CCAGGCATCA TTAACAAGC CAGTAGGTCA 180

35

CCTGGCTTCC GNGGGCCAAT TCATCTTTCA GACAATCTTT AGCAGAAATG GACTCAGGGA 240

AGAGACTCAC ATGCTTTGGT TAGTATCTGT GTTCCGGTG GGTGTAATAG GGGNTTAGCC 300

CCNGAAGGNN CTGAGCTAAA CAGTGTTATT ATGGGAAAGG AATGGCATTN CTNGTTTCAA 360

CNAGCGCTAN ATGCAACCAT TNCNCTNTTG TTATAGTATC TAAGGGTNNG NNAGTTAAAC 420

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GGTTN 425

SEQ ID NO:6916

SEQUENCE LENGTH:81

SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS08147

SEQUENCE DESCRIPTION:

GATCCACCAC CCAGCTTCAC CCCTCACCCC CAGCGGCTCA CCATGGGGAT GGCAGCAATA 60

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AAATACTTCT AATGGAACAA A 81

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SEQ ID NO:6917
 SEQUENCE LENGTH:254
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08148
 SEQUENCE DESCRIPTION:
 GATCAACAAG CTTNCCACG AATTAAAGAG TTTTATCAAG ATATATCGAA TACTTCTACC 60
 CATCTGTTCA TAGTTTATGG ACTGATGTTT CAAGNTTGTA TCATTCCCTT GCATATAATT 120
 AAACCTGGAA CAACATGCAC TAGNTTATG TCAGNAATAT CTGTTGGNTT TCCAAAGGNT 180
 GTTAACAGAT GANGTTTATG TGCAAAAAG GGTAAGATAT AAATNCAAGG AAGNAAAANN 240
 GTTGNTAGCT AAAN 254

SEQ ID NO:6918
 SEQUENCE LENGTH:134
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08149
 SEQUENCE DESCRIPTION:
 GATCAGAGGA ACCTTAGAGG CCTGAAATTG TTGCTTCCAG TTTAGCTGCC CCTCAAATTC 60
 AAGTGAATAT TTTCCCTTCT CCCTTACCC TTCTCCAGAA ATAAAGCAGG TGACAGGGTT 120
 TTNAGAATCT TAAA 134

SEQ ID NO:6919
 SEQUENCE LENGTH:277
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08150
 SEQUENCE DESCRIPTION:
 GATCAAGAGT AATTACCAAC TTAATGTTTT TGCATTGGAC TTTGAGTTAA GNTTATTTTT 60
 NAAATCCTGA GGACTAGCAT TAATTGACAG CTGACCCAGG TGCTACACAG AAGTGGCTTC 120
 AGTGAATCTA GGAAGACAGC AGCAGACAGG ATTCCAGGAA CCAGTGTTTG NTGNGGCTAG 180
 GNCTGAGGAG CAAGCGAGCA AGCAGAGNAA GAAAAGTTAA ATACCAGNTA AGCTTTTGAT 240
 TTTTGTATTG TTTGCATCCC CTTGCCCTCA ATAAACN 277

SEQ ID NO:6920
 SEQUENCE LENGTH:255
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08151
 SEQUENCE DESCRIPTION:
 GATCTACCAC CAGAGAGTTC ACACCTGNGA AAGNCCTCAT GAGTGCAATN AATGTGAAAA 60
 ATCCTTTAGC CGAAGCTCCA GCCTCATTCA CCACCGGAGA CTTCACTG GAGAAAGACC 120
 CTATCGGTGC AGTAAATGTG GGCAGTCATT TAAGCAAAGC TCCAGCTTCA GTTCACATCG 180
 GAAAGTCCAC ACNNGGGANN GGCCTTATGT GTGTGGGGNC TNTGGGNANT CCTTAGCCA 240
 TAGTCCANC CTTAN 255

EP 0 679 716 A1

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SEQ ID NO:6921
SEQUENCE LENGTH:273
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08153
SEQUENCE DESCRIPTION:
GATCTGGCTT NTCCNGGAGG CTNCCATGGT TGGAAGATGG TATCAGAGGG CCTGCCTGGG 60
CAGTCTGTCT CCGGGCCAGG GTCAGGGACC CTCTGCCTCT GGCAGCCTTA ACCTGTCCTC 120
TGCTAGGACC AGGGTGATT CAAGCCAGGG AAGCAACTGG GACCCTGAAA ACTGTCCCTC 180
CCCAGCCCGG TCCNCGTGTA TGTGCCNTGG TCCCCTTGCT GCCATGTGGA TGCTGTTGTN 240
ATTGCTGTTT GTATATTATC AAAATGTTTT TAN 273

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SEQ ID NO:6922
SEQUENCE LENGTH:83
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08154
SEQUENCE DESCRIPTION:
GATCTTCTTT ATAATATTT ATTGTTTGAA TCACTTTTAG GATGTAACCT TATAAATAAA 60
CATGAGCGCT GATGATTGC AAA 83

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SEQ ID NO:6923
SEQUENCE LENGTH:207
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08155
SEQUENCE DESCRIPTION:
GATCTCATC TCCATCCTCA TCCTCTTCT TCACAGCATT TACTTGGAGC TCTTTGTGAC 60
ACACCATGTC AGTCATGATG AATCGGCCAA CAGCCAGCCC TTGCCAGCTG ACGTCACAGT 120
CTAAGATGGG AAAGTGTGGT ACAGATAGAC ATGAAGAGAG CTTAGCAGTG ATTGAGGTGG 180
TGAATAAATA TACAGTCATT GAATAAA 207

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SEQ ID NO:6924
SEQUENCE LENGTH:199
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08156
SEQUENCE DESCRIPTION:
GATCACTGTG GGCTGGAGAG GAGAAGGAAA GGGTCTGCGC CANNCTGTCC GTCTTCACCC 60
ATANCAAAGC AACTAGAGC AAAAAACCAG TTGTAATATA AAATGCACTG CCCTACTGTT 120
GGTATGACTA CCGTTACCTA CTNITGTCAT TGTATTACA GCTATGGCCA CTATTATTAA 180
AGAGCTGTGT AACATCAA 199

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SEQ ID NO:6925

EP 0 679 716 A1

SEQUENCE LENGTH:358
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 5 CLONE:HUMGS08157
 SEQUENCE DESCRIPTION:
 GATCATGCAG CAGAAGCAGA AAAAGGCAAA CNAGAAGAAG GAGGAACCCA AGTAGCTTTG 60
 TGGCTTCGTG TCCAACCTC TTNCCCTTCG CCTGTNTGCC TGGAGCCAGT CCCACCACGC 120
 10 TCGCNTTTCC TCCTGTAGTG CTCACAGGTC CCAGCACCGA TGGCATTCCC TTTNCCCTGA 180
 GTCTGCAGCG GGTCCCTTTT GTNCTTCCTT CCCCTCAGGT AGCCTCTCTC CCCCTGGGCC 240
 ACTCCCGGGG GTGAGGGGGT TACCCNTTCC CAGTGTTTTT AATTCCTGTG GGGCTCACCC 300
 CAAAGTATTA AAAGTAGCTT TGTAATTCCA AAAAAAANGG NNNNNAGGGN GGGGGGGN 358

15 SEQ ID NO:6926
 SEQUENCE LENGTH:62
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08158
 20 SEQUENCE DESCRIPTION:
 GATCTGTACA TTTTACTATC TGTAATAAT TCTTCAAAGA ATAAATAAA AAAAAAGCA 60
 AA 62

25 SEQ ID NO:6927
 SEQUENCE LENGTH:51
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08159
 30 SEQUENCE DESCRIPTION:
 GATCAGATAC AGCATTAAAC GTTCTCAGT CTCCTNTCC TCATCTGTAA A 51

35 SEQ ID NO:6928
 SEQUENCE LENGTH:259
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08160
 SEQUENCE DESCRIPTION:
 40 GATCTTGCTA AAGTTAAAT AAGGAACATT TCACCTTTTA AATATTTAAT TCTNATGTGG 60
 ACTTATTTCC AGAAACTTT GGTGATAATT CTTGAGACAA AAGGTGGTTA AGTAGCATT 120
 TTATGTAATG CTTATATACC ATAGAGTTTT TAATAGANGA GAAATCCATT TCCTCCGAGG 180
 GTCACTATTA ACAATGTACT TCCTTAAAT TNGTTAATG ATTGTGATGG GTGCTGCATT 240
 45 TGCACATTGC ATTANGGAN 259

50 SEQ ID NO:6929
 SEQUENCE LENGTH:226
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08161

SEQUENCE DESCRIPTION:

5 GATCGGATGC TGAATAAAAC TCACCGTGAA GCAAGTCCCA CTGAACGAAA AGTTGGTCGT 60
AGCATTNTC AGCAGCTGCT TCCGGCTCCC TGAGACCCAG CCATGGAAGG CATAGAAAAG 120
CATGTGACAA GAAACTCATG GCTTTTAATC CTGGCAGGNC AAGGAAAACC CCGTCAAAGN 180
TTTGATAAN TTTTGCTGCT CTCCTGCTGG GGGGGGNNNG NNNNNN 226

SEQ ID NO:6930

10

SEQUENCE LENGTH:219

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08162

15

SEQUENCE DESCRIPTION:

GATCGTTCTT GATTTTGTTT TCATTAGTGT CATTTCCTTG TCATTGAGGA CTTTCCCT 60
TACAACAGTA ACACCATTTT TTGAAGAGCA AAACCTATAA TACCTCCTGG GATTGTNAGC 120
TAGTCATTCA GCCTGTGTAA CCATGTGGAA ATAAAAATTG ACGACCAATG TATTATATGG 180
ACAACTTTTG CTTTGAGTAA TAACTTGAT TGTAGGAAA 219

20

SEQ ID NO:6931

SEQUENCE LENGTH:111

SEQUENCE TYPE:nucleic acid

25

TOPOLOGY:linear

CLONE:HUMGS08163

SEQUENCE DESCRIPTION:

GATCCACCCC ATTCTTTCAC TTTAAGAAAA AACAAATAAT TGTTCAGAG GTGTCTGTAT 60
TTTGCAGCTG CCCTTTTGTA AGANGCACTT TTCCCAAATA AAACAATTAA A 111

30

SEQ ID NO:6932

SEQUENCE LENGTH:109

SEQUENCE TYPE:nucleic acid

35

TOPOLOGY:linear

CLONE:HUMGS08164

SEQUENCE DESCRIPTION:

GATCATTGA AATNTCTGGG AGTCTGAGGN GTACTGACAT AATNACCTGC TGGAGTCTGT 60
AAATACACAT TTAAGACAGT GAGGATGTGA ATAAATATAT TAATGCAAA 109

40

SEQ ID NO:6933

SEQUENCE LENGTH:111

SEQUENCE TYPE:nucleic acid

45

TOPOLOGY:linear

CLONE:HUMGS08165

SEQUENCE DESCRIPTION:

GATCAAATAG AGCCCTGTCC AGGTAGAAAA GAAATGGTAT GTAGAGCTTA GATGTCCCTA 60
TTGTGACAGA GCCATGGTGT GTTTGTNNTA ATAAANCCAA AGAAACATAA A 111

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EP 0 679 716 A1

5 SEQ ID NO:6934
SEQUENCE LENGTH:243
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08166
SEQUENCE DESCRIPTION:
GATCCCAGCT GTTCTAGGCA GGGCCAGGCA GAGTGGGGCG CAGGCCCTNG AAGGGCGAGA 60
10 CCCAGTGGCT GGGCTGCCCC GGGCTGAGGG GCCGCCTCTT GAGGGTACAC GCCTCTGGTC 120
ACATGGCCAT GGAGCCTTGG GTACCCCTGA GTTAAGGGAG GACATTGGC CAGCTGGTGG 180
CTGGGAGGGG AGCCTGGCTG CCCTGCTGCT TCTCCTGCCT AATAAACAGG CTTCTCCTGC 240
AAA 243

15 SEQ ID NO:6935
SEQUENCE LENGTH:124
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08167
20 SEQUENCE DESCRIPTION:
GATCTATTTT NATAGGTTCT TTTTCAGAAG TAAAATTTTG TACATATATA CATGTACATA 60
TCTNTTTAGT TTGGGTTTCAT TTCTATAACA TTTTGTAGA AAATAAAAGT TTGAGCACCT 120
GAAA 124

25 SEQ ID NO:6936
SEQUENCE LENGTH:169
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
30 CLONE:HUMGS08168
SEQUENCE DESCRIPTION:
GATCAGTTGT ATGTAAATGT ACATTTTTGT NACTTTGGCT GTGCCCGTTA GAATTTATCT 60
TCCATAAAGT ATTTCTCCCA TTGAGTCTAA TGATGTATAC TTTGCCTAGG TCTTTCCAAA 120
35 ATTAAATTTA TGTAATGTC TATTTTATAT AAAATATGNT TAAAATAAA 169

40 SEQ ID NO:6937
SEQUENCE LENGTH:314
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08169
SEQUENCE DESCRIPTION:
GATCACATTT AGGAGCAAGC AGGTGAGGGC AGAGGTGCTG GAGTGAACGT CGGCTCTTTG 60
45 GCTGGTAGCC TTACCCACTG AGGGAGAGGA GCTGAAACAT GAAATGGGGA TCGGTGANCA 120
ATGTCTCCCT TTTGCAGAGC TGTATTGACT TGTCAAGAA TAACTGATAT GCCTTCACTC 180
AGAAGAAAAG AAATGAATGT GAAAGAAAGC CAAGCATCNN TTTGCACTTA AATCANTTAC 240
CACGGAAGNT ATATTTAGCT TCAACTTTAG TTTAAAAATTA TGTGAATTAA ATATTTTGAT 300
50 TTCCCTGGGN NAAA 314

55

5 SEQ ID NO:6938
 SEQUENCE LENGTH:46
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08170
 SEQUENCE DESCRIPTION:
 GATCAAGTGA CTGTTTAATA AATNAGTTGC TTGGGTAGAA TGGAAA 46

10 SEQ ID NO:6939
 SEQUENCE LENGTH:21
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS08172
 SEQUENCE DESCRIPTION:
 GATCTTGCTG AAAACCAGAA A 21

20 SEQ ID NO:6940
 SEQUENCE LENGTH:85
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08173
 25 SEQUENCE DESCRIPTION:
 GATCAACAAT CATCACCTGC CTTTGTAGA AAAGAAAAAC AAAAAAGTA AATAAAATT 60
 TTAAACAGTA AAATAAAAGT TTAAG 85

30 SEQ ID NO:6941
 SEQUENCE LENGTH:274
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08174
 35 SEQUENCE DESCRIPTION:
 GATCACTACT GCAGCAGAAT GGTCTATGTN ATTTTCCTTC TTTGGTTTTT NCCTGACTTA 60
 CATTCGTAAT TTTAAGAAAA TTTCTTTACG GGTGGAAGCC AATTACATG GATTAACCCT 120
 CTATGACACT GCACCTTGCC CTATTAACAA TGAACGAACA CGGCTACTTT CCAGAGATAT 180
 TTNATGAAAG GATAAAATAT TTCTGTAATG ATTATGATTC TCAGGGATTG GGGNAAGGTT 240
 40 CACAGNAGTT GCTTATTCTN CTCTGTAATT TTAN 274

45 SEQ ID NO:6942
 SEQUENCE LENGTH:246
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08175
 SEQUENCE DESCRIPTION:
 50 GATCTAAGTT GGGCAAAAGA GAGTCATCCT GTTACCTTAA AAAGGAGTCT AGGGTGAGCC 60
 CCAGTCTGCT GGTGGCCATC TTCACCACCT TTGGGGAGAG ATGGCCTCNG CAGACACACA 120

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EP 0 679 716 A1

5 AAGCAAAGAG GTGGAGAGAA ACTGTGTCTT TGTGTCATTG TTTGAGTCCC TGGACCCAGC 180
CCAGTCCCCC CGACTACGCT GANCTTTTCA ATTGTGGGGG GCAATAAATT CCATTAGTT 240
ANGAAA 246

10 SEQ ID NO:6943
SEQUENCE LENGTH:215
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08176
SEQUENCE DESCRIPTION:
GATCCACATN ATACCGGTGC TGAAGACCTG CAAGTNATTT TGGAAAAGGG CTGGTGCGGA 60
TGGAAGGGCC CAGATTTTGG GAACAAGGAT GCATACTATA ACTTATGTCT TCCTCAGCGA 120
15 CCAAATATGA TTAAAAATAT CTTGGAGTCA AAGACTGCAG TAGAGTGGTA TTATAAATTT 180
NTGAATAAAG AATCAGTTTA NTTTTTCACA TTAAT 215

20 SEQ ID NO:6944
SEQUENCE LENGTH:171
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08177
SEQUENCE DESCRIPTION:
25 GATCTGGACA CGAGCAGTCC CTGAGGGGGCG GGGTCCCTGG CGTGAGGCCC CCGTNACAGC 60
CCACCCTGGG GTGGGTATTG TGGGCACTGC TGCTCTGCTA GGGAGAAGCC TGTGTGGGGC 120
ACACCTCTTC AAGGGAGCGT GAACTTTATA AATAAATCAG TTCTGTTTAA A 171

30 SEQ ID NO:6945
SEQUENCE LENGTH:161
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08178
35 SEQUENCE DESCRIPTION:
GATCGGGCCC GGAAAAAGTC TGCCCCAGAA ACTCTCACAC TTCCAGACCC TGAGAAAAAA 60
GCCACCCTGA ATTTACCCGG CATGCACTCT TCAGATAAGC CATGTCGGCC CAAATCTGAG 120
TAACTTTATA TAANTATCTC ATGGGGTTTT ATATTTTCAA A 161

40 SEQ ID NO:6946
SEQUENCE LENGTH:251
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
45 CLONE:HUMGS08179
SEQUENCE DESCRIPTION:
GATCCTTGGG GAAGGTTTCA GTTGCACTGT ATGCTGTTGG ATTTGCCAAG TCTTTGTATA 60
ACATAATCAT GTTTCCAAAG CACTTCTGGT GACACTTGTC ATCCAGTGTT AGTTTGCAAG 120
TAATTTGCTT TCTGAGATAG AATATCTGGC AGAAGTGTGA AACTGTATTG CATGCTGCGG 180
50 CCTGTGCAAG GAACACTTCC ACATGTGAGT TTTACACAAC AACAAATGAA AATAAATTTT 240

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AATTTTATAA A

251

SEQ ID NO:6947

SEQUENCE LENGTH:277

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08180

SEQUENCE DESCRIPTION:

10 GATCCCTCTT GGTGTGAGTT TTCTGACCCA ACCAGCCTCT GGTAGCATC ATTTGTACAT 60
 TTAAACCTGT AAATAGTTGT TACAAAGCAA AGAGATTATT TATTTCCATC CAAAGCTCTT 120
 TTGAACACCC CCCNCACTTT ANTCCNTNGT TCAGGCCGAT GNGCTTGCTT TCCTTCANCC 180
 TGTGTGTTTT CTTATTTAGG NCTATTTATT ANTGGTTGGN CCAATGTACT CACAGCTGTT 240
 15 GCGTCGAGCA GTCCTTNGTG AAANTNCTGT ATAANTN 277

SEQ ID NO:6948

SEQUENCE LENGTH:130

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08181

SEQUENCE DESCRIPTION:

20 GATCCAGCTT CAGCTGCACA CCTNTGTCCC CTTGGATGGG GNACTAAGGG NAAACGTCTG 60
 25 TTGTATCACT GAAGTTTTTT GTTTTGTTC NATACGTGTC TGCATAAAAA TGCCAAAGTT 120
 TTTTTTCAAA 130

SEQ ID NO:6949

SEQUENCE LENGTH:101

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08182

SEQUENCE DESCRIPTION:

30 GATCACCAAT NNTNNCTTTG AGCCAGCCAA CCAGATGGNG AAATATGACC CTCGCCATGG 60
 35 TAAATACATG GCTTGCTGCC TGTAAATACCG TGGTGACGTG N 101

SEQ ID NO:6950

SEQUENCE LENGTH:304

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08184

SEQUENCE DESCRIPTION:

40 GATCATGAAG ACCTGCTTTA GCCCAACAG AGTGATTGGA CTCTCAAGTG ACTTGCAGCA 60
 45 AGTAGGAGGG GCATCAGCTC GCATCCAGGA TGCCCTGAGT ACAGTGTTC AATATGCAGA 120
 GGATGTACTG TCTGGAAAGG TGCAGCTGA CAATACTGTG GGCCGCTTCC TGATGACCT 180
 GGTTAACCAA GTACCGAAAA TAGTTCCCGA TGACTTTNAG ACCATGCTCA ACAGCAACAT 240
 50 CAATGACCTT TTGATGGTGA CCTACCTGGC CAACCTCACA CAGTCACAGA TTGCACTCAA 300
 TGAN 304

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EP 0 679 716 A1

5 SEQ ID NO:6951
SEQUENCE LENGTH:268
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08185
SEQUENCE DESCRIPTION:
GATCCTATCT GTNCTGGCAG GACTGGCAGN CATGGTGGGC TACGCCTTGC TCAGNGGCAT 60
10 TGTCTCCATN CAGNGGGCAA CGCTGCTCGG GCGCCAGGCA CCCGGACCCT GGGCATGGCT 120
GAGGAGGATG AAGAGGAATN ATTTGTCCTC ACGCTCCCAA GACTGGTTTT TNTACTCTNA 180
TGCAATCCAG NGGCCCCCGT GCCTCCTCGT TGTGGTACA GCGGGACACG GGGTGCTGCC 240
ACCCAGAATA AAGCCACTCA CACTGAAA 268

15 SEQ ID NO:6952
SEQUENCE LENGTH:152
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08186
20 SEQUENCE DESCRIPTION:
GATCCCACCG AAGCAGCGTG TCCTCCGTGG GAGCCCGAGG TGGGTCCAC CCGCATCTCC 60
TCTTCCACAC CCTCCTGGGA CCCCAGCCCT GGAGAAGTAG CTCTGGCTTG TTTCTTTTT 120
NATTAAAGAT AATTTATGTA TCGACTAAGA AA 152

25 SEQ ID NO:6953
SEQUENCE LENGTH:72
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
30 CLONE:HUMGS08188
SEQUENCE DESCRIPTION:
GATCAGTGGG TCTAAGTNTC CGAGACTTAA CGAAAATAGT ATTCAGCTG CAATAAAGAT 60
TGAGTTTGCA AA 72

35 SEQ ID NO:6954
SEQUENCE LENGTH:293
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
40 CLONE:HUMGS08189
SEQUENCE DESCRIPTION:
GATCAGCATC CTNCGNTGCC CCTTAGCAAC TNAGGTGGTT GATTTGAAAC TGTGAAGGTG 60
TGATTTTTTC AGGAGCTGGA AGTCTTAGAA AAGCCTNGTA AATGCCTATA TTGTGGGCTT 120
45 TTAACGTATT TAAGGGACCA CTTAAGACGA GATTAGATGG GCTCTTCTGG ATTTGTNCCT 180
CATTGTGCAC AGGTGTCTTG TGATTGANAA TCATGAGCGA AGTGAAATTG CNTTGANTTT 240
CAAGGGAATT TAGTATGTNA ATCGTGCCTT NGAAACANGG GGGNAAAAAN ATN 293

50 SEQ ID NO:6955

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SEQUENCE LENGTH:184

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08190

SEQUENCE DESCRIPTION:

GATCCTTCAA TAAAAATCA ATGTGTGAAG GTTCCTAGGC ATAAAAGGAA GTTCAGAAGG 60
 NTATGTAGGA AATACTGTTG TCTCTCTGGG GATTGGGATT ACAGGGATTT TTTCTCTTTT 120
 NCTTTTGTGTT AAGTTTGTGT TTCCTAAAGT TTCTGCAATA AACGTACGAT GCTTGATATAA 180
 TAAA 184

SEQ ID NO:6956

SEQUENCE LENGTH:327

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08191

SEQUENCE DESCRIPTION:

GATCTTGAAT TTCTACATAA CTTTCTCAGT TTATATGCCC TGTGGCAAGT GCAGCAAGCA 60
 CTGTTTCTG TTTCTAAACT TGTAGAAAAT CATCCATACA TCTTACAGTT GTCAGTTTTA 120
 ACCAGATAAC AGTGGCACTT TGTGCTGCT TTTTATCTT TAGCTTAGGT TAACAGGACC 180
 CTGGAAGTAA AGTTGTTGAT TTATTCAATA GAGTATTCTC AATTAATTG GCTAGATTTC 240
 TACATGATTC AAANTCTAAA AANGTAGACA TGCATGCTTA CATGTCTAAG GCCTGAAAAA 300
 TTGGNGGGGN CATCCCAAAA TAAATGN 327

SEQ ID NO:6957

SEQUENCE LENGTH:22

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08192

SEQUENCE DESCRIPTION:

GATCTAAGTG GTGGTATGGA AA 22

SEQ ID NO:6958

SEQUENCE LENGTH:223

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08193

SEQUENCE DESCRIPTION:

GATCTGTTGA CAATGGAGCA AGTGAAAGAC TTTGCTGCTA ATGTGTATGA AGCTTTTAGT 60
 ACCCCTCAGC AACTGGAGAA ATGATTTTCC CTTCAAGAAA AACTACAGTG GGATTCAATT 120
 ACTTTTAAA ATACACTGGG TAAATCACCT ATACTTAGAG TAACAGTTTG TTATCAAAAT 180
 GCCTGNTAAA ATATATNCTT AATAAANGTC TTCATTTCAT AAA 223

SEQ ID NO:6959

SEQUENCE LENGTH:331

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08194
 SEQUENCE DESCRIPTION:
 5 GATCTTACTT GCGAAATGCG ATGGTTGCTG GGAATACCTG AACTGTGGA TTATATTGCT 60
 TGACTTCTAC CTCAGAATCT NCTTTGTTTC ATGACTTAAT AGTGCTTTAA GTNNGGTATA 120
 TNATTTNACC NCTAGGAATT CTNTGTTTCA CACAGAAATA AAAATTTTAA AATAGAAAAT 180
 GCTTTTACTT TGTAAGGTAA GNGAGTATCC ATATGCTTAG ATGTGCTCGT TTCTAAAAATN 240
 10 CTAGNGGTTG CTATANTCAG CTCATGGAAT GCACAGCTAT TGCTTTTGT GNTAGATTGT 300
 ACATAACAT CAGCAGNTGN AAGGGNAAAT N 331

SEQ ID NO:6960
 SEQUENCE LENGTH:211
 15 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08195
 SEQUENCE DESCRIPTION:
 20 GATCACCCAG ATGCAGTGAA GCTTTTNACC CGCCAGCTTT TGGAAGTCCA TCAGGGACAA 60
 GGCCTAGATA TTTACTGGAG GGCTAATTAC ACTTGTCCTCA CTGAAGAAGA ATATAAAGCT 120
 ATGGTGCTGC AGAAAACAGG TGGACTGTTT GGATTAGCAG TAGGTCTCAT GCAGTTGTTC 180
 TCTGATTACA AAGAAGNTT AAAACCGCAA A 211

25 SEQ ID NO:6961
 SEQUENCE LENGTH:208
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08196
 30 SEQUENCE DESCRIPTION:
 GATCATTGTA GATGAAGTGA AGCAAGAAGT TATCAGTACC AGCAGCAAGG CAGANCCNCC 60
 CCAGTGCACC TCCCTGGCCT GGTCTGNTGA TGGCCAGACT CTGTTTGCTG GCTACACGGA 120
 CAACCTGGTG CGAGTGTGGC AGGTGACCAT TGGCACACGC TCGAAGTTTN TGGCAGAGCT 180
 35 TTNCAACTAA ANAANAANCT GGCATAAA 208

SEQ ID NO:6962
 SEQUENCE LENGTH:165
 SEQUENCE TYPE:nucleic acid
 40 TOPOLOGY:linear
 CLONE:HUMGS08197
 SEQUENCE DESCRIPTION:
 GATCCCAGCC CTAGGTGGG CAGAGGCAGA CCCTCCCAG AGCTCCTTAG GAAGAAGACA 60
 45 GACTGGTTCA TTGAATGCCG CTTATTTAT TTCTGGTGAG GATGCATGCG TGGGGCTGCT 120
 GGTGTTTAGA GTGGGGGCTA CCCTTNAAA TCACTGCTAC TCAA 165

50 SEQ ID NO:6963
 SEQUENCE LENGTH:280
 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08198
 SEQUENCE DESCRIPTION:
 5 GATCACTCTN GGTCAAGCTN TCTTTCTTTA TCTNTCTTTC TCCTTTTTTA AGAAAACGAG 60
 TTAAGTTTAA CAGTTTTGCA TTACAGGCTT GTGATTCATG CTTACTGTAA AGTGGAAAGTT 120
 GANCTTATTT TAAAACTTCA AGCTCAGTAA TTTTGAACAC TGNAACATTC ATCTAGGNCA 180
 TAATAACAAA GTTCAGTATT GACCATAACT GTTAAACAA TTTTAAGCTT TCCTCAAGTT 240
 10 AGTTNTGTTG TAGGNGTGTA CCTAAGCAGG TAAGCGTANA 280

SEQ ID NO:6964
 SEQUENCE LENGTH:51
 SEQUENCE TYPE:nucleic acid
 15 TOPOLOGY:linear
 CLONE:HUMGS08199
 SEQUENCE DESCRIPTION:
 GATCCATATA TGTGTGCATG TCATGAAATA AAAGNATCAC ACANCACAAA A 51

20 SEQ ID NO:6965
 SEQUENCE LENGTH:155
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08200
 SEQUENCE DESCRIPTION:
 GATCTTGGAG CTTCCCTGTA GCCCACCTTC CCCGTGCTTC ATGTTTGTAG AGGAACCTTG 60
 TGCCGGCCAG GCCCAGTTTC CTTGTGTGAT ACACATAATGT ATTTGCTTTT TTTGGAAATA 120
 30 GAGAAAATCA ATAAATTGCT AGTGTTTCTT TGAAA 155

SEQ ID NO:6966
 SEQUENCE LENGTH:58
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS08201
 SEQUENCE DESCRIPTION:
 GATCCAGCAT TGGGGCCACT AACTCCAGC CTGGACCACG GAGCGAGACT GTCTCAA 58

40 SEQ ID NO:6967
 SEQUENCE LENGTH:111
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS08202
 SEQUENCE DESCRIPTION:
 GATCAGGCTT CAGGATAGCC CAGCACTGAG GACATCTAAA ACAGTCTTCA TTTATTTATT 60
 TATTGTTTAT AGTGTCCAAA AGTTGGAATT AAAGTGAATG GCTGAATCAA A 111

50 SEQ ID NO:6968

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EP 0 679 716 A1

SEQUENCE LENGTH:39
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08203
 SEQUENCE DESCRIPTION:
 GATCATGACA AAGAAGATTA AAATTCATT AGCATGAAA 39

SEQ ID NO:6969
 SEQUENCE LENGTH:90
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08204
 SEQUENCE DESCRIPTION:
 GATCAATTGT ACCCCATACC TCAGCATCAT GCAATATAAC CATTAAAGAGA CCTGCACATG 60
 TATTCCCTGA ATTTAAAATA AAAGTTGAAA 90

SEQ ID NO:6970
 SEQUENCE LENGTH:315
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08205
 SEQUENCE DESCRIPTION:
 GATCATTGCA GTCTCTGGCT TTTNACTCC AGGCGGGCTC CGTGCCCTCC GGGGCCTCTT 60
 CTTCTTTGTN AGGCCAGTGG GAGCCACAGA GGCCTCATTG GCTTCCTATG AATATNAAAT 120
 GGGATTCTCT GTGGGGGAAG CTTCCATAAG GCAGGAAGGG GTCCCCGGTT GTTGGCTATN 180
 AGTGTATTAT NCTTTTAAAT CATCTGTGCT CTTTCAAAGA CGACGTAAGA AGCTTGTGCC 240
 CTGGCGTTGC AGAAATCTCT GGCTGTAGGG ATTAAACCTA CATGTCTTCC TGATGGGGTG 300
 GTGAGGACGT AAACN 315

SEQ ID NO:6971
 SEQUENCE LENGTH:262
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08206
 SEQUENCE DESCRIPTION:
 GATCACCTGA GGTGAGGAGT TCACGACCAG CCTGGCCAAC ATGGTGAAAC CCTTCTCTAC 60
 TGAATATACA AAAATTAGCC AGGTGTGGTG GCGCATACCT GTAGTCCCAG CTAATTGGGA 120
 GGCTGNGGCA GGGAATCAC TTGAACTCGG GAGGCGAAGG TTGCAGTGAG CCGAGATTTC 180
 ACCAGTGCAC TCCAGCCTGG GTGACAGAGC AAGACTCCAT CTCAAAAAAA ANNTAATTAN 240
 AAAAAATTGC AGCTGCNGGA AA 262

SEQ ID NO:6972
 SEQUENCE LENGTH:118
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS08207

SEQUENCE DESCRIPTION:

5 GATCTTGAG GTCTAGTATT TNAATAATGC ACTATTACCC AGGGCAGNTA TTATGAGAAA 60
CTGTTTCTNC TCTAAGGGTT TATGGCAGAC TTTGCTTTT NAACATGTGA GANATGAN 118

SEQ ID NO:6973

SEQUENCE LENGTH:152

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08208

SEQUENCE DESCRIPTION:

15 GATCTGAGTA TTCCCAAGG GCACTCAGCC ACAGGGNTGA ACTAGTGTTC GTTTGTATCC 60
TTGCAGTTAT GTGACTTAAA GTGACTCAGA GATATCTATG TATGCNNGTG TACTCCGCTG 120
GCAATGAAGT AAATATATCA ATTTGTGGNA AA 152

SEQ ID NO:6974

SEQUENCE LENGTH:297

20 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08209

SEQUENCE DESCRIPTION:

25 GATCCTCCTG CTTTGGCCTC CCAAAATGCT GGGNTTATAG GCATGAGCCG CTGCACTTGG 60
CCTGNTACTG ATTTTAATNN CTGCGTTAT CACATAGTGT TGTATTGAA ACATAGTTCA 120
TGGTTTATC AAAGAACTGA AGNTGAGAAT ACTGGTCATC TAACTTTGTA ATTTGATTG 180
ATTATACTGT AAAGTTTGAC AGTCTCATTT TATCACTGCG TTTGTATCTA TNACTAAANT 240
GTATTTNTTG ACCTCTTACT GATTCATTGT TGGTNTGTNC AAACGTGGA CTNGNTN 297

SEQ ID NO:6975

SEQUENCE LENGTH:301

SEQUENCE TYPE:nucleic acid

35 TOPOLOGY:linear

CLONE:HUMGS08210

SEQUENCE DESCRIPTION:

40 GATCCTGTCA CCCAGGTAGT GAGTATAGCA CCCAGTGAAA CTGTAGTCTC ATNCCAGGCA 60
CTGTGCTAGC CCACTCTGGG TCATTTAATC CTCTCCTAAG AAGAGAGGAG ACACAGCGTC 120
CCCATTGAC AGATGCAGAA AGAGGTTCCA CAGGTGTGCN TTGATTCTNT CCTAAAACCG 180
TTTCCCGGAA GCTTTTCCTG GTGTGGGCGC TTCTAACCTA ATCCTCAATC GNTTCCAGNA 240
CTATTACTCT GTTCCACAG TGNTACTGTG TCTAGGTTTT ANGGGCGGCA AGTTCNATTT 300
N 301

45 SEQ ID NO:6976

SEQUENCE LENGTH:264

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

50 CLONE:HUMGS08211

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SEQUENCE DESCRIPTION:

GATCCTTCCG ACTTGGCCTC CCAAAGTGTT GGGATTACAG GCATTAGCTA CCACACCTGN 60
 CCAAGGCCCA GGTTCGACA GAAAGGGAGA GAAACCTGC CAGAGATGCC ATTCGGAGC 120
 CACTCTGCTT GGCAGGGACC TGTGTTCCCC TCATGCAGGT TCATCCTTAG AGGGCTGCGG 180
 TCTTATCTGG TTGTGCAAAA GTCCACAAC CTTTCTGGAT TGATAGTTTG TGGTGAANNT 240
 AAACAATTTT AGTTTGTGTTG GAAA 264

SEQ ID NO:6977

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08212

SEQUENCE DESCRIPTION:

GATCAGAACT ACAATTACAA CCAGCATGCG TATCCCACTG CCTATGGTGG GAAGTACTCA 60
 GTCAAGACCC CTGCAAAGGG GGGAGTCTCA CCTTCTTCCT CGGCTTCCCG GGTGCAACCT 120
 GGCCTGATGN AGTGGGTGAA GTTTTGGTAG GCAATTTCTT GCAACCACCA CCGNGGCCCC 180
 GNAAGCACT GGTGCTNAGG GAGCTCCTNN CCTTGGCCCN CACGCCTGTG CCAGCCCTGG 240
 NCCCGNCTGC CACACCTCTG TTTCTTANGG CTGGGGACNC AGATTGTNTC TCCTTNTTTC 300
 TTNCNACTGC AATGTGCN 318

SEQ ID NO:6978

SEQUENCE LENGTH:166

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08213

SEQUENCE DESCRIPTION:

GATCATNCTG CCCATNTCC ACGCACCCAA AACCAGCCCT CTCGCTTCTA ACACAGGGCA 60
 CTTGCTGGGG CTCAGGGATG TTAGGGACGA GTTCCAGNCC TGCCACTNCC CTNGGGCGAC 120
 CCCTCNCCTG CCCTGCCTCC CTGCTCTGCC GCCCCTNTTN CTGGAN 166

SEQ ID NO:6979

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08214

SEQUENCE DESCRIPTION:

GATCCGATAC AGATTACATT TCTTGCCAGA GAGTGTGCT GTTGTTCCTT TAGGTAAGTG 60
 TGTATTGGTG ATTCCATAAT AAAAGCAGTA ACAATAACTA CCAA 105

SEQ ID NO:6980

SEQUENCE LENGTH:233

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08215

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

5 GATCAGATAC ATGAACATGT CTTACATGGG TTGCTGTATT TAGAATTATA AACATTTTTC 60
ATTATTGGAA AGTGTAACGN GGACCTTCTG CATACTGTG TAGAACCAAA ACCACCATGA 120
CACAGTTTTT ATAGTGTCTG TATATTTGTN ATGCAATGGT CTTGTAAAGG TTTTAAATGN 180
AAACTACCAT TAGCCAGTCT TTCTTACTGA CANTAAATTA TTAATAAANT AAA 233

SEQ ID NO:6981
SEQUENCE LENGTH:72
SEQUENCE TYPE:nucleic acid
10 TOPOLOGY:linear
CLONE:HUMGS08216
SEQUENCE DESCRIPTION:
GATCTTTCCA GAGGTCCATG GTGGAAGACG ATAACCCTGT GAAATACTTT ATAAAATNTC 60
15 TTAATGTTCA AA 72

SEQ ID NO:6982
SEQUENCE LENGTH:58
SEQUENCE TYPE:nucleic acid
20 TOPOLOGY:linear
CLONE:HUMGS08217
SEQUENCE DESCRIPTION:
GATCACCCAT CTGTGTGCTT CCATCCTGCA TTAATAATCA CTCAGTGTGG CCCAGAAA 58

25 SEQ ID NO:6983
SEQUENCE LENGTH:280
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
30 CLONE:HUMGS08218
SEQUENCE DESCRIPTION:
GATCTCCAAC AGCCAGTGTG TGTTCCTCAT CTCTTGTAGG TTCCATCAAT GGTGAGCACC 60
AGCCTGAATG CAGAAGCGCT CCAGTATCTC CAAGGGTACC TTCAGGCAGC CAGTGTGACA 120
35 CTGCTTTAAA CTGCATTTT CTAATGGGCT AAACCCAGAT GGTTCCTAG GAAATCACAG 180
GCTTCTGAGC ACAGCTGCAT TAAANCAAAG GAAGTTCTCC TTTTGAACCT GTCACGGATT 240
CCATCTTGTA AAGGATATTA AATGTTGCTT TAACCTGAAA 280

40 SEQ ID NO:6984
SEQUENCE LENGTH:311
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08219
45 SEQUENCE DESCRIPTION:
GATCCAGAGA CTCGTGCTGT CTCTTTGTCC CTTTCCTGAT GGTGTTACTT CGATTAACTA 60
CTTCAGCTTG GTGTAAAAAT TGCCAAAGCA GTTGCCCTGCC ACAACTTTGT AAAAGCCAAA 120
AAGGAGGTTG AAAATTCACA GGCTGCCCGA AAAAAGANGA AACTTGCATG GGGGTTTGAA 180
GCAAAGNNGN GATGGGAAAC CNAAGCAAC ATGGGNTACA TGTAACCTGC CAGAGTGCTT 240
50 CAAGACATTT GTAGCCTCAA ATGCTCAAAT TTANTGNAA TGTTCCTG CCTATGTTCA 300

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TATGTCAGGT N

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SEQ ID NO:6985

SEQUENCE LENGTH:160

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08220

SEQUENCE DESCRIPTION:

GATCTGATGG TTTTATAAAG GGGNGTTCTG CTGCATACAC TCTGTCTTGC CTGCTGNCAT 60
 GCAAGATGTG ACTTAGCTCC TCCTTTACCT TCCACCATGN TTGTNAGGCC TCCCAGCCA 120
 TGTGGAAGCTG TGAGTTGANT AAACCTCTTT CCTTTATAAA 160

SEQ ID NO:6986

SEQUENCE LENGTH:115

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08221

SEQUENCE DESCRIPTION:

GATCAAAAGA TAACTGCTTA TNATAGTAAG ATAACCTTGG AACAGAACTG TGATTAAAAA 60
 CCTGAATAAT AAAAAACACA CAAGACCCAT TAAACCTAT TCCTCACTTT TAAAA 115

SEQ ID NO:6987

SEQUENCE LENGTH:331

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08222

SEQUENCE DESCRIPTION:

GATCCAGTTG GATATAGTAA CCACTGCCGA CTNCTGGAT GTGCTAACTC ACACCAAGCC 60
 CTCCGNAAG AATCTGGCTC AGAGATACTC AGACTGGCAA AGAGAATTCG AGTCTGTCTG 120
 AAACCACATT TACCCTGACC TGGCCACAAA GGCAACCACA AAGACCTCCT AGTTTATTAN 180
 NGNCGTGGG AGAACAAAAT GATTGGAATG GAAAAGAGAA AATTATTTT GANGACTGGA 240
 TTAACCTGAG CCACTGTATT GTTTTGNTA GCTGAGATAT ATTTNTTAAC TTTACCATTA 300
 TCGATGTCAG CAAAATATTG AGAGTTTCTT N 331

SEQ ID NO:6988

SEQUENCE LENGTH:121

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08223

SEQUENCE DESCRIPTION:

GATCTACTCC CAGTTTTTAA CTTTTNACAG AATATTGTTG CTTTCCTACA ACAATGTACA 60
 TATATTTNCA GTTGATATG CTTTTTTTTT TNCCAAATAA AGTTTGAGTT CTAAACTCAA 120
 A 121

SEQ ID NO:6989

EP 0 679 716 A1

SEQUENCE LENGTH:151
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 5 CLONE:HUMGS08224
 SEQUENCE DESCRIPTION:
 GATCCTTCCC TGGCTCTCAC CTGCTCCTTT AGGCTGTAAT TGCTTCCTAC ACACGAAGCC 60
 TCTGATTGGA GCCTCTGGTN CATCTCAGAA AAACCTTCCA AGAGCGCTGG GGTTCATGCT 120
 10 TTCTGAATAA ACACTACTGT TTACATGGAA A 151

SEQ ID NO:6990
 SEQUENCE LENGTH:119
 SEQUENCE TYPE:nucleic acid
 15 TOPOLOGY:linear
 CLONE:HUMGS08225
 SEQUENCE DESCRIPTION:
 GATCTCCATA ATTTTAAGAT ACTCTATGTT AAACCTTTT TNAAGTNCTC TTTTACATC 60
 20 ACGTCTGAAA TGCACGAGNG TGGCGGTTTC TGTTTCACTG GTTTNNTTGT TCATTTTNN 119

SEQ ID NO:6991
 SEQUENCE LENGTH:116
 SEQUENCE TYPE:nucleic acid
 25 TOPOLOGY:linear
 CLONE:HUMGS08226
 SEQUENCE DESCRIPTION:
 GATCCCGGGC CGTTATCCAT CTGGAGGCTG CAGGGTCCTT GGGGTAACAG GGACCACAGA 60
 30 CCCCTACCA CTCACAGATT CCTCACACTG GGGAAATAAA GCCATTTCAG AGGAAA 116

SEQ ID NO:6992
 SEQUENCE LENGTH:207
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS08227
 SEQUENCE DESCRIPTION:
 GATCCGCTGG GACAGAGTGC CCACCGTCGT GTCTTTNATA TCTCTGTAAT AACCCACAGT 60
 40 TAAAAAGGC AGGAAAAGGC ATCAGGGTAC CATGTTATCA GGCAAGTTCT TACATTCTAA 120
 GAAAATTCCT AGCTTTGTTG GGTGAGGCAA GTTTTAAAN GAANAATGTC ACTTCCCCAT 180
 TTGATAANGA TTTCGTTATC ANTTAAA 207

SEQ ID NO:6993
 SEQUENCE LENGTH:325
 SEQUENCE TYPE:nucleic acid
 45 TOPOLOGY:linear
 CLONE:HUMGS08228
 SEQUENCE DESCRIPTION:
 50 GATCCTGATT GTAAGGTTTT CCACCTATTT CCCTGGATAT TTTNNTGGTC AGTACTGGCT 60

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EP 0 679 716 A1

5 CTGGTGGGTG TTCCTTGTTT TAGGCTTTCT CCTGTTTCTN AGAGGATTTA TCAATTATGC 120
 AAAAGTTCGG AAGATGCCAG AAACCTTCTC AAATCTCCCC AGGACCAGAG TTCTCTTTAT 180
 TTATTAAAGA TGTTTCTGG CAAAGGCCCTT CCTGCATTTA TGAATTNTGG TCTCAAGCAG 240
 CAAGNGAACA CCTGCAGGAN GTGAATCAAG CTGCAGAACA CAGAGGNNTA CTCACCTGCT 300
 TTAATAAAT ANAGTACTGT TGATN 325

10 SEQ ID NO:6994
 SEQUENCE LENGTH:130
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08229
 SEQUENCE DESCRIPTION:
 15 GATCCCTCGG GNNTTAANTA CANGAAACCT AGAACTGACC AAAAGGCATT ATAACCTCTGA 60
 CTCAAATACA AGGTACAGAA GATAAGCATC TTTGAGGAAA CTCCTACTTC AGTTCTTTTG 120
 TTATNATGNN 130

20 SEQ ID NO:6995
 SEQUENCE LENGTH:116
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08230
 25 SEQUENCE DESCRIPTION:
 GATCATATAA CATTCCAGGT ACTCAGGATG AATGGTTTGA GGAAGGTCT GAATCTTCA 60
 AAGGTTTCAG CTGTATTAAC ATTCTCCATC TAATAAACTT TATCTTGTC TGAAGA 116

30 SEQ ID NO:6996
 SEQUENCE LENGTH:33
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08232
 35 SEQUENCE DESCRIPTION:
 GATCAGCAAA TAAATGAAAT TTATTAAG AAA 33

40 SEQ ID NO:6997
 SEQUENCE LENGTH:246
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08233
 SEQUENCE DESCRIPTION:
 45 GATCTTTAGG CAGTTTTTAC TTGGCCAGAA AGCAGTGCTG AATACTTGNA ACTGTGTGCT 60
 CTGTTCTACT TAATGTTCTG TCAGAATGTT CTTTGTAGG CAGTATGTCA TGNTGTAATC 120
 ATCTATCTCC TNGTCTGTTT CCAAGTTNNN CTGTGAAGTC TGCGACCCTT TTGNGGTGGT 180
 CATCANAGTC ACAGATTCCT TGTTTAACCA NGTGTCCNA AGCATGTACC TGNAGTTATA 240
 TCATNN 246

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5 SEQ ID NO:6998
 SEQUENCE LENGTH:186
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08234
 SEQUENCE DESCRIPTION:
 10 GATCATTATG CAATAAGCT GCTAAAGTCA CATTGTGGT TGATACTCAA GCCTTGTTACA 60
 AGCACATATC ATGTAGACAT TTTAAATGT GAAGCTTTAA CAGGNATAGG ATAGCTGTTT 120
 CAAACCTTGT GCAATTGTAA ATAANTTGT TAGCTGTGTT TCTTTCATT TTAATANAAA 180
 NTTAAA 186

15 SEQ ID NO:6999
 SEQUENCE LENGTH:154
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08235
 SEQUENCE DESCRIPTION:
 20 GATCTTCATA TTAATCTTAA AATTTTGTA CGTGTCTTT TCCTTTTTT CCACAGTTT 60
 AATATATNAT TCITCAACGA CATTTTTGT AACTTTACAC TTTTGGTT ATNTATTTT 120
 AAAAAATGA AAAATTAATT TAAAAAATG CAAA 154

25 SEQ ID NO:7000
 SEQUENCE LENGTH:195
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08236
 SEQUENCE DESCRIPTION:
 30 GATCTGGTAC GNGGNGGACG NCATGGCCCG AGATGCCAC GGAACACAG CGCTGACCTA 60
 CGCCCGNAG GCCTCCAGCC AGGAGTGCAT CAACGTGCTT CTGCAGTACG GCTGCCCCGA 120
 CGAGTGGGTG TAGTATCTGT TTTATTNNC TGNAGTCTCC TTGGTGTAAC AACAAAATNG 180
 35 GNAAAATAAG GATAN 195

40 SEQ ID NO:7001
 SEQUENCE LENGTH:163
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08237
 SEQUENCE DESCRIPTION:
 45 GATCAGGCC TAAACGTTG TTGTGATGAG GTTCTTTAG CAAGTTCTTG TTAAATTAT 60
 CATTTATTTG ATGAGTGAAG TTTTAAACAT GCTTTGCTGT GTGAAATTA AAAAGGGAT 120
 GTTTTCCAG GCTGGAACAA TAAATGTGC TGTGCAGTT AAA 163

50 SEQ ID NO:7002
 SEQUENCE LENGTH:127
 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08238
 SEQUENCE DESCRIPTION:
 5 GATCTTTGTG AAAGTAGTAC AGTATATNAC CTTTAATTNC TTTTTATTT TAAATATACT 60
 GTNACCNTGA AGCACTGGTT GGGCATTITA ATTCATGTTA ATAAATCACA ATTATGTCAG 120
 TTTTAAA 127

10 SEQ ID NO:7003
 SEQUENCE LENGTH:123
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08239
 15 SEQUENCE DESCRIPTION:
 GATCAGCCTC CTGTCTTATG TGGGCACGTT CCAAAGTTTA AATGCATTTT TTGACTCTT 60
 GGCCAAAATT TAGAAGATGC TGTGAATATC ATTTTGAAC TGTGTAAATA CATGAAAGAG 120
 AAA 123

20 SEQ ID NO:7004
 SEQUENCE LENGTH:260
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08240
 SEQUENCE DESCRIPTION:
 GATCTCAGCA TTACCTCTTT GGNAAAGGAA GGTAGTTCAA GAAATGAAGA GCTGTTGATG 60
 GGATGATTGA AGAAACAGCT ATGAGAGGAT TGGCTCCCAT CTTTGTNAC TCTTGGGNCA 120
 TCCTGTCTATC TGAGAATGAA CAAAGACCAA TTTTGTGT GTGAAGCTTA AGGGTCATAT 180
 30 GTTGCTTGT NTTTNTNAAT GCTAATCCTT GTGAAANTAA ATTGNCCAGG CGGAAGGAAA 240
 NCTCTATTTT AGGATNCAAA 260

35 SEQ ID NO:7005
 SEQUENCE LENGTH:79
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08241
 SEQUENCE DESCRIPTION:
 40 GATCTGTATT TNTATATACA CGTGTCCGNA CAAGTATAAC TAAATAAAAA TTAAAGNTTT 60
 TAATCATTTT AATTGGAAA 79

45 SEQ ID NO:7006
 SEQUENCE LENGTH:59
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08242
 SEQUENCE DESCRIPTION:
 50 GATCATATTT CTGCCAAGNA TGTGCCTTCA ACTTTATAAT TATAGTGTG TAAAATAAA 59

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EP 0 679 716 A1

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SEQ ID NO:7007
SEQUENCE LENGTH:67
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08243
SEQUENCE DESCRIPTION:
GATCTCACTT TTNCTTATTA GAATTTGTAC CTTTGTGTTGA TTAAGGAAAT AAAAGATGAA 60
TGACAAA 67

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SEQ ID NO:7008
SEQUENCE LENGTH:76
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08244
SEQUENCE DESCRIPTION:
GATCAGTCTC TCAGTGCCTN CCTACTGGGC AGCTCATCTG TCCACTTATT CGTATTAAAT 60
TTNCTTTTTA TTAAAA 76

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SEQ ID NO:7009
SEQUENCE LENGTH:162
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08245
SEQUENCE DESCRIPTION:
GATCCACATC CCANATGCTG AACAGAGAG CCCCTCATCC CTCCTCCCAG TACTTAGGCA 60
GCGCCTTCTG TGTGCACTGC CTCTTGCCGG GCACTGGGTA GGTGCACACG NGGACCGGGA 120
CAGGGCCAGA NACTTTGGTN CCCGNGGTTT GTNTGNGGGG NN 162

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SEQ ID NO:7010
SEQUENCE LENGTH:58
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08246
SEQUENCE DESCRIPTION:
GATCGATTAA GTCTAGGAGT TTNAGACTTG CCTGGGCTTA GCAAGACCCT GTGGCAAA 58

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SEQ ID NO:7011
SEQUENCE LENGTH:275
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08247
SEQUENCE DESCRIPTION:
GATCCTTGAG TCTCCNCTCT GGATGGAATC CGCGAGCTGG CCACCTGGCC ACCCTCTACA 60
CGACTCCACC CNCCATGGC CGTGGGGCCC TTACTCTCTG ACTTCTCAGG ACACAGGTCA 120

EP 0 679 716 A1

TGGAGGTTCT TCCCAAGCTG GCAGAGGCCA TTTGTGAAA GTGGAGAGCT ACGTGGTGGN 180
CATCTGCCAA CTCCAGCATC TCTGGNGGGG NTCCACGNTG AATGTNATTT TTGAAAACAG 240
CTTATGTANT TAAAGGTTGA ATGGCACATC ATAAA 275

SEQ ID NO:7012

SEQUENCE LENGTH:124

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08248

SEQUENCE DESCRIPTION:

GATCAAGACC ACCCTGGCTA ACACGGTGAA ACCCTGTCTC TACTAAAAAT AAAAAAANTT 60
AGCCGGGCAT GGTGGCACAC GCCTGTACAT CCCAGCTACT CANGGNGGCT NGTGGCATGG 120
NGNN 124

SEQ ID NO:7013

SEQUENCE LENGTH:215

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08249

SEQUENCE DESCRIPTION:

GATCANGAGC CCTTTNCTC CAGTAGTACA GGCTTGAAA ACTACTTCTA TTAAGTTATT 60
GATGCAATTT GATATTTTT CATAATCTAT ATTTAAACAA AATTACATCA TTGCATCATC 120
TTTTCTAAAT TCATCTCCAT TAAAACTGC CTTAAGCTAC CAGNTTGCTT TTGCCANNGN 180
NGGCCATACT GTGTGTTGT TTGTTAATN NNCTN 215

SEQ ID NO:7014

SEQUENCE LENGTH:373

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08250

SEQUENCE DESCRIPTION:

GATCTCTCAG GGATAGAAGA TATTCTCAT GAAGGCAGCC TAACTCTGAG GAAAACAATG 60
CCAATTCAAG TACAGATTTC AACACATCTT CAACACTATG TGAAGGGTTC ACATCTTAAC 120
CTGTGCAATT CAGATTGATA CTCAGAATAT GGGTTGATTT GAATATCTGA AATATCAATG 180
GAAAATCCCA CTNNGTTTT GATGAACAGT TTGAACAGTT TTCTGTAATC AAGCAGCTTG 240
CATAGAAATT GTATGATGAA ATTTNCGNN GGTCTTGGT GCTGTTTTGT TCTTTTTTTG 300
TTTTTTGTTG TTTTGTATT TACTTATATA CATATAANCT TTTNTTGAAA ATATGTTTTG 360
GTTACTAANA GTN 373

SEQ ID NO:7015

SEQUENCE LENGTH:219

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08251

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

5 GATCCCAACC TGGACCCACT CATGAACCCT CACATCCGCG TGGGCCCCTC CTNAGCCCN 60
 TTGCTTGTGG CTAGGCCAGC CTAGGATGTG GNTTCTGTGG AGGAGAGGCG GGGTAATGGG 120
 GAGGCTGAGG GCACCTNTNC ACTGCCCTC TNCNTCAAGC CTAAGNCACT AAGACCCAG 180
 ACCNAAAGCC AAGTCCACCA GAGTGGCTGC AGGNCAGGN 219

10 SEQ ID NO:7016
 SEQUENCE LENGTH:91
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08252
 SEQUENCE DESCRIPTION:
 15 GATCGCTTGA ACCATAAGC TGGAGACCAG CCTGTGCACA TGGTGAGACC CTGTCCTAC 60
 CAAAAAAT AAAGTCATTA GAGTTTTTAA A 91

20 SEQ ID NO:7017
 SEQUENCE LENGTH:176
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08253
 SEQUENCE DESCRIPTION:
 25 GATCTGATGG TTTATAAGG GGCTTCCCCC TTCACTCAGC TCTCATTNTT CTCTCTCCTG 60
 CCACCATGTA AAGAAGGATG TGTGTTGGTTT CCCTTCCATC ACGATTGTAA GTTTCCTGAG 120
 GCCTCCTAAG CCATGTGGAA CTGTGAGTCA NTTAAATCTC TTTACTNTAT NNNNGN 176

30 SEQ ID NO:7018
 SEQUENCE LENGTH:174
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08254
 SEQUENCE DESCRIPTION:
 35 GATCATCACT TACCCACCCC CCAAGTTCAA GACCAAATCT TCCAGCTGCC CNNTTCGTGT 60
 TTCCCTGTGT TTGCTGTAGC TGGGCATGTC TCCAGGAACC AAGAAGCCCT CAGCCTGGTG 120
 TAGTCTCCCT GACCCTNTN AATTCCTTAA GTCTAAAGAT GATGAACCTC TAAA 174

40 SEQ ID NO:7019
 SEQUENCE LENGTH:45
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08255
 45 SEQUENCE DESCRIPTION:
 GATCCTCCCT TTCCTATAGN GNTAAAGTG ATTTATCTTG GCAAA 45

50 SEQ ID NO:7020
 SEQUENCE LENGTH:109
 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08256
 SEQUENCE DESCRIPTION:
 5 GATCATGTGA TGTTGGAGTA GGAGGGTTTA TNTTGGTGAC TTTGGGATGC TGGTTTCATT 60
 GTAGGTATAA TNATGNGGNG CAAAGAATCC AGGNAAGAAT TGCCAGAGN 109

SEQ ID NO:7021
 10 SEQUENCE LENGTH:225
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08257
 SEQUENCE DESCRIPTION:
 15 GATCCCACTG AGGCCAGAGA ATAGGGTCTG GAGACAAAGG AGCATTCACT TCAGCCTCTG 60
 ACTGGTGGCA GGCCAAGTNT TTATTACAT AGGGTGTAA CAAATAGGAA ACCTCTAAAG 120
 GGTACTTAAA CCCAGATTT TNTACACAGG GCACTTGCTT GAGCCTCATC CCGCTTTCTG 180
 GAATGTACTT TTGCTTCAAT AAATCTGTGC TTTTGTTCCT TCAA 225

20 SEQ ID NO:7022
 SEQUENCE LENGTH:196
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08258
 SEQUENCE DESCRIPTION:
 GATCTGGAGG GCGTGGCTAC AGGACCCGGG ATGCCATTCA GTTACTCATC TTTNATGCTT 60
 TCGTCCTGAC CTGTCTCAAC TAGACTTGCT CCTGCAACCA CCATGGGGGT TTTGCATTTA 120
 30 CATTTGTGGA CCATGTTACA GTTAAGAAAA ATCCTGTTTC AGTCCTTATA TGTAATAAAA 180
 TGTTTTATGA TGTAAA 196

SEQ ID NO:7023
 35 SEQUENCE LENGTH:170
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08259
 SEQUENCE DESCRIPTION:
 40 GATCTNTCCT CCGCTTGAGA AGACCATGGA ACTGGATAGC TCACTTCCGG AATACAACAT 60
 GTTCCCCGC AAAGGCCGGG GNCTCCCTAG NTCAAATTC CAATCTCATN AGCAGGTGNT 120
 AGAAAGGTNT TGTGGTAGCA ACTNCATGAT TTCTCTGTGA CAAAAATNAN 170

45 SEQ ID NO:7024
 SEQUENCE LENGTH:84
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08260
 50 SEQUENCE DESCRIPTION:
 GATCAAATGT AAATNTTTTG AGGATATTTT NCTTGTTTTT NAAAATCAAT TTTTAACCAA 60

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ATATTAAACC CTTTATTTC CAAA

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SEQ ID NO:7025

SEQUENCE LENGTH:246

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08261

SEQUENCE DESCRIPTION:

GATCCACCAC TGGNCTCTCC CTCCCCCAGC CTGGAGCACG GGAGGGGAGG TGACGGCTGG 60
 TGA CTGATGG ATGGGTAGTG GGCTGAGAAG AGGGGACTAG GAAGGGCTAT TCCAGGCTCA 120
 GCCCTGCTAC TGCAGCTTTG CCGCTGAGTG TAGGAAAAAC AGGCATGACA GACCAGGGTG 180
 AGGGTTGTGC CCAGCTGGGC CACGGCCATG CGTGGGGTGG CCCAATAAAC ACCGTGGACT 240
 CCCAAA 246

SEQ ID NO:7026

SEQUENCE LENGTH:45

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08262

SEQUENCE DESCRIPTION:

GATCAGTCTT TATNAATAAA ACCTGTTCTC TTTAATCAGC TTA AAA 45

SEQ ID NO:7027

SEQUENCE LENGTH:186

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08263

SEQUENCE DESCRIPTION:

GATCAGACTG NTAACCTTAA CACCATAGGT GCAAAAANGAG TTGTTCTNAG GTTTTTTTGA 60
 AACATTAAAG TTCCAAAACA TGACATTTTT AAGAATAAAT TTGAAATAGN GTATGATTGA 120
 ATGCAGAGNA TTATGTACCT CTAATTGCTT AATTTTGTAG NGGTCTTNA TTGTAGAATT 180
 GGNNCN 186

SEQ ID NO:7028

SEQUENCE LENGTH:271

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08264

SEQUENCE DESCRIPTION:

GATCTCTTGG TAACAATTGA CATTGCACTA GNATTTGAAT GNTACTGTTG TTGGCATTTC 60
 TGTTANTCTT TCAAGTAGTT GTAAATAATC TCTTTTCTC ATATAAACGG AATGCATAAT 120
 GTCTTACTCT TGAGTCATTT AGTAGCATTG ATATGGTACC AAGTTGTATA GCAGTGTAAC 180
 AAAATCATGC TTTGACAATG NAATTTACAT ATTGCCTCAC CTAGAGATGT TTTGTCTGTA 240
 TGACTNNTNN CNTTNTAAAG TTNGNCNAAC N 271

EP 0 679 716 A1

SEQ ID NO:7029
 SEQUENCE LENGTH:301
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08265
 SEQUENCE DESCRIPTION:
 GATCAATCA GGGTAATTGG GTATCCATCA CCTCAAACAT TTACCATTTC TTTGTGTTGG 60
 GAACATTCCA TATCTTTTTT TCTAGCTGTT CTGAAATTG CTATACGTTT TTTAACTAA 120
 AGGTCCTTG CTGTGTTATG GAACACTGGA ATTTGTTCTT TCTGTTTAA TGTATTTTGG 180
 TACCTATTAA CCAACTTCTC TTTATCCTCC CCCGTCCTCT ACCCTTCCCC AGCCTGTGGG 240
 AATCACCCT NCTACTCTCT GCCTTTATGA GATTTACTTN TTTNATCTCC CNATATAANA 300
 N 301

SEQ ID NO:7030
 SEQUENCE LENGTH:117
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08266
 SEQUENCE DESCRIPTION:
 GATCGTAGAA ACACAGAATG GGAAGTGGGA AGCCCTTTGG AAATCCAGCT GCAGAAACAG 60
 ACACCCCAAT GCTATTTACA TACAGCTCTA TATATATAAA AAAAGNAAAT ATGNAAA 117

SEQ ID NO:7031
 SEQUENCE LENGTH:297
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08267
 SEQUENCE DESCRIPTION:
 GATCTCACCA GGGCCCAGTC ATATATGCAC AGTTAGACCA CTCCGGCGGA CATCACAGTG 60
 ACAAGATTAA CAAGTCAGAG TCTGTGGTGT ATGCGGATAT CCGAAAGANT TAAGNGAATA 120
 CCTAGAACAT ATCCTCAGCA AGAAACAAAA CCAAACTGGA CTCTCGTGCA GAAAATGTAG 180
 CCCNTTACCA CATGTAGCCT TGGNGACCCA GGCAAGGACA AGTACACGTG TACTCACAGA 240
 GGGAGNGNAA GNTGTGTACA AAGGNTATGT ATAANTNTTC TTTTAGTCA TCCTATN 297

SEQ ID NO:7032
 SEQUENCE LENGTH:58
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08268
 SEQUENCE DESCRIPTION:
 GATCCTGGGG ACTCATATTC TTTCAGAATC ATGTAAATAA ATGGCATCAT GTTGTAAT 58

SEQ ID NO:7033
 SEQUENCE LENGTH:103
 SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08269
 SEQUENCE DESCRIPTION:
 5 GATCTATTG GCCTCTCAAA TNACTNAGA TTCCTGTAA AAAAGATTGA TGTTATTGTC 60
 TCTTGTAGAG GAAACTAATA AAGTGTGTGT ACCTNTGTGN AAA 103

SEQ ID NO:7034
 SEQUENCE LENGTH:265
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08270
 SEQUENCE DESCRIPTION:
 15 GATCAAGAGT NATGGGCCTT GGAGCCCTTG CCAGCCCAGC CCACCTGTNA GGAGGCTGCC 60
 CATCCCCTCT TTNAGGCCAC CCTGTGTCCT CTCCTGCTGCC TCTGCCCAGA GCTCCAGCCG 120
 GAGTGTCTTG CTGCTCAGAC CCCTCCGAGG TCCAAGTCCT GTCCTGCCAC TCAGCTCCCN 180
 CTGCACCCCA GAAGAAACGC AGGGTGCGGT TGCATTTGAT TTCAGATAAA CAACAACCTC 240
 TTAGTAAAAA NACCTNCCCA CTAAC 265

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 SEQ ID NO:7035
 SEQUENCE LENGTH:46
 SEQUENCE TYPE:nucleic acid
 25 TOPOLOGY:linear
 CLONE:HUMGS08271
 SEQUENCE DESCRIPTION:
 GATCAATCTT GANGTGAAAT ATTTCCAAAA TAAAATTCTA CAGAAA 46

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 SEQ ID NO:7036
 SEQUENCE LENGTH:254
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS08272
 SEQUENCE DESCRIPTION:
 GATCTTTGCA TAGAATGTCA AGCTAACCAG GCGTCCGNTA CTCAGAAGA GTGTACTGTC 60
 GCATGGGGAG TCTGTAACCA TGCTTTTAC TTCCACTGCA TCTCTCGCTG GCTCAAAACA 120
 CGACAGGTGT GTCCATTGGA CACAGAGAG TGGGAATTCC AAAAGTATGG GCACTAGGAA 180
 40 AAGNCTTCTT CCATCAGGNG TTAATTGTTT TGTATTTCAT TTAATGGCTT TCCCTGCTGT 240
 TACCTAATTA CATN 254

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 SEQ ID NO:7037
 SEQUENCE LENGTH:222
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08273
 SEQUENCE DESCRIPTION:
 50 GATCATTGAG AAAGTGTGTTG AAACCTTCTC ATGAAGTGTA TATATAATGG CGTGAAAAAT 60

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EP 0 679 716 A1

5 TCCTTTGGAA AAATNTATGT NCCTTTCATT TTGCCCAAAT TGCAAATTTT CAGCATGGNT 120
 GTGAAAAGCA TTAATAATTAT AACTTTGTGT ACANGATGAA AATAATTCAC TAAATTTGCC 180
 CTTTTTTACA CAAAATAAAA TGTAAANGTT ANNGCTGGNA AA 222

10 SEQ ID NO:7038
 SEQUENCE LENGTH:32
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08274
 SEQUENCE DESCRIPTION:
 GATCAAAATGA CANTAAATCA TTCAAAATCA AA 32

15 SEQ ID NO:7039
 SEQUENCE LENGTH:109
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08275
 20 SEQUENCE DESCRIPTION:
 GATCACCCAG GTACANCATG CTCTATATCC ATGCGGCCCT AGAAACCCAA GGAACATATA 60
 CAACCACTCT TGTGTCCCTC CTTATAACCN CAGNTTTATN CNAGAAGTN 109

25 SEQ ID NO:7040
 SEQUENCE LENGTH:301
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08276
 30 SEQUENCE DESCRIPTION:
 GATCACAAAT TGGTGCATTC TTAAGTGTGG ATAATTATGT ATGAAGCCAA ATTGTAATAA 60
 AACTGTTATT TGCCAAACAT AGAANCCTTG CTTGCTGACT AGGCTGAGCC CAGAGTGTTA 120
 TTCTCTTCTT CTGCCTGTNA TGTATCTGGG CAATTTGTAG GGCCTTTGCA GGTGAATCCA 180
 35 GTAGAGAACC ATGCTAAGAN TGNGCAGCAG TCATGCATGC CTGTATCATT GAAAGCCTTC 240
 CTNGCCATTT TNNCAGCTTC ACCTTNCGTG GTTTTTATTT TTTCCAGAGG GGGCCTGGGG 300
 N 301

40 SEQ ID NO:7041
 SEQUENCE LENGTH:151
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08277
 45 SEQUENCE DESCRIPTION:
 GATCCTCCTC CCTAAGCGCT TATCTTGGTG CTTCCAGCTC AGTCTGGCCA ATTCCACCTC 60
 TATGAGATAA GGAGGGTTCA GAGAACCCAG TCACTTTGCT GTTTTCTGC NCACAGAGTT 120
 TCTCTAATAA ATCTTATTCT ATGTCTACAA A 151

50 SEQ ID NO:7042

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SEQUENCE LENGTH:138

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08278

SEQUENCE DESCRIPTION:

GATCACATTG TTTGACCCAG TATGTCTTGT AGACACGTTA GTTATAATCA CCTTGTATCT 60

CTAAATATGG TGTGATATGA ACCAGTCCAT TCACATTGGA AAAACTGATG GTTTTAAATA 120

ANCTAATTCA CTAATAAA 138

SEQ ID NO:7043

SEQUENCE LENGTH:300

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08279

SEQUENCE DESCRIPTION:

GATCTGTCAA CAGGTTAAGT CAATCTGGGG CTTCCACTGC CTGCATTCCA GTCCCCAGAG 60

CTTGGTGGTC CCGAAACGGG AAGTNCATAT TGGGGCATGG TGGCCTCCGT GAGCAAATNG 120

TGTCTTGGGC AATCTNAGGC CAGGNCAGAT GTTGCCCCAC CCACTGGAGA TGGTGCTGAG 180

GGAGGTGGGT GGGGCCTTCT GGGGAAGGTGA GTGGAGAGGG GCACCTGCCC CCCGCCCTCC 240

CCATCCCCTA CTCCCCTGN TCAGCGCGGG NCATTANAAG GGTGCCANAN ATTTTTTTNN 300

SEQ ID NO:7044

SEQUENCE LENGTH:90

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08280

SEQUENCE DESCRIPTION:

GATCTCATTC CATGGGAAAA AAAAAATCC TGTCTTGTTC ATAAATTGAC AATGTCACTA 60

AATAGAAATA AAGTGCACAN TAAATGNAAA 90

SEQ ID NO:7045

SEQUENCE LENGTH:175

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08281

SEQUENCE DESCRIPTION:

GATCTNAGTT AGACATCACC ANTGGAAATC GATTGGAATA GAAACTTAAA GGAAATGGAA 60

CCCTGACTAT TCTCCCATCA AATCATATAT GTTGACCTGT CTGAATTATA AACCAGCCTG 120

ACCTTTCCTT TAGCATTAGA TGTAAATAAA TAACCTTGGGA AATTTGTNAT TTAAA 175

SEQ ID NO:7046

SEQUENCE LENGTH:190

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08282

EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCCTCAGG ACTGACCACT TAGCTGGCAG GTTGTNCAGC TTTTAATTCC AGTCATAATA 60
GGTGACAGTG TTAACCGTGA AANTTTGNGA GGCACNTTTG NCCTCTTCCC TATAAAAATC 120
ACACAAGCGT GNTTTTACAA AGGGTCCCCG TGGGAACCTT NGCTTAAGGA CCCTTTGNCC 180
CNTCAGTTAN 190

SEQ ID NO:7047

SEQUENCE LENGTH:147

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08283

SEQUENCE DESCRIPTION:

GATCCTCAGG ACTGACCANT TAAGCTGGCA GGTGTGCCA GCTTTTAAAT CCCAGTCATA 60
ATAAGGTGAC CAGTTTAAAC CCGTGNAAC TTNAGGGGA CTTTGCCTT TTCCCTATNA 120
AATNACACNG GGGGGTTTAA NAAGGTN 147

SEQ ID NO:7048

SEQUENCE LENGTH:223

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08284

SEQUENCE DESCRIPTION:

GATCCATCTG CCCTCATAAA AGTGTTCAGG TACAGCAGCT GAGGCTGCCC TGAGNAATCA 60
AGGGGCCATT ACCAAGGGGC AGGAAAAGNA TATGTAAGAG GTGGCCTTCA TGGTAGAGCT 120
TGACCCAAGA ACTACTCCAN ATTCGGATGG CCCAGACTGG GNGCATCCCC TGACTTTCCC 180
TTTGACTTCA CCCTGTTTGT AAATAAANCA AGGNAGATGC AAA 223

SEQ ID NO:7049

SEQUENCE LENGTH:56

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08285

SEQUENCE DESCRIPTION:

GATCTCCTCC CTNGTTTAAA TGTCAATAAA TGCCCCAACT GCTTTGTAAG TGCAAA 56

SEQ ID NO:7050

SEQUENCE LENGTH:107

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08286

SEQUENCE DESCRIPTION:

GATCACCTTA AAATTATTN ACGNACTGNA GNCAATNAAG TCCGTTATGT TTAGAGTAGA 60
AAATGTTTGA GTTAAAGAGC ATCTGTCAAC AGAATCTACA AAAAAGN 107

SEQ ID NO:7051

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08287

SEQUENCE DESCRIPTION:

GATCCAATTN CATNNGCGGC TGCATCTNCT GCTCATGCCG CCCAAACACC ACCCAGATGT 60
CACTTACGTC AAGAAGGTCC GGACCCTCCG TATGCACCTG TTCACGGCCC TGCAGCTGCT 120
NTGNN 125

SEQ ID NO:7052

SEQUENCE LENGTH:276

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08288

SEQUENCE DESCRIPTION:

GATCCAAGAA ACGNANGACA CAACAACAAC GATGCCGTGT GTCTGCTCTA CCTGGGCAAG 60
CTCAAGNACT CCTGCGGCA GCTNGAGGCC ATGGTCCAGC AGGACCCAG GCACTACCTG 120
CACGAGAGCG TGCTCTTCAA CCTGACCACC ATGTACGAGC TGGAGTCCTC ACGGNGCATG 180
CAGAAGAAAC AGGCCCTGCT GNGGGCTGTC GNCGGCAAGG NGGGGGCCAG CTTAACACA 240
CAGTGCCTCA AGCTGGCCTA GCTGNNTNNN ACACAN 276

SEQ ID NO:7053

SEQUENCE LENGTH:209

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08289

SEQUENCE DESCRIPTION:

GATCGAGTTT ACAGTTGTGN NNTCTTGAAG GTATTACTTA ACTTCACTAC AGCTTGTCTA 60
GAAGACCTTT CTAGNGTTA TCTGGTTCTA GAAGGGTCTA TACTTGTCTT TGTCTTTAAG 120
CTATTTGACA ACTCTACGNG TTGTAGNAAA CTGCTAATAA TACAAATNCT TGTGTCCAT 180
GGNAAGGCAA ATAAATTTNC TACAGTGNN 209

SEQ ID NO:7054

SEQUENCE LENGTH:126

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08290

SEQUENCE DESCRIPTION:

GATCTCATTT TCAGTTTAAG TAACTGCTGT TACTTAAGTG ATTACATTT TGCTCAAATT 60
GAAGCTTAAT GGAATTATNG CCCTCAGGAT AGTATTTTGT AAATAAAGNT GATTTAAATA 120
TGNAAT 126

SEQ ID NO:7055

SEQUENCE LENGTH:276

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08291

SEQUENCE DESCRIPTION:

5 GATCGCACCA TTGCACTCCC ACCTGGGCAA CAAAGAGTGA AACTTGGTCT CAGAAACGAA 60
 ACAAACACA AAAACCTTTC TNAGTCCCAG CATATGTGGA GCAGCCTCAT TCTTCATAGC 120
 TGTGTGTCAT TCCGTTGCGT GATGGGGTCA CAGAGCACAG ACCTGGTGCC CTTTTCCTTT 180
 TTAATATGTG GAAACCCCTC CATGCTTTCC AAAGCCTACA AGTACAGCAG CCCCAGTTT 240
 10 AGGGTGNGCA GCAGTGGTCA GAGCTCTTN CTANTN 276

SEQ ID NO:7056

SEQUENCE LENGTH:276

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08292

SEQUENCE DESCRIPTION:

20 GATCTCCAGG AGTCCCCTGC ACATACCTTC TCCATCGTGT CAGCTGTGTT TCTCTTGATT 60
 CCGTGACACC CGGTTTATTA GTTCAAAAGT GTGACACCTT TTCTGGGCAA GGAACAGCCC 120
 CTTTAAGGAG CAAATCACTT CTGNNNNAGT TATTATGGTA ATATGAGGCA ATCTGATTAG 180
 CTTACAGAC TGAGTCTCCA CAACACCAAA ATATCCAGAT GTAAACCCCA AACTTGTACA 240
 CAAANGNAAG CACAGATTGT TTACCTGTTG TGGAGN 276

25 SEQ ID NO:7057

SEQUENCE LENGTH:167

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

30 CLONE:HUMGS08293

SEQUENCE DESCRIPTION:

GATCCCTGTT ACCCAGAGAA TANNTACATT CTTTATCTTG ACATTCAAGG CATTTCATC 60
 ACATATTGA TAGTTGGTGT TCAAAAAAAC ACTAGTTTGG TGCCAGCCGT GATGCTCAGG 120
 CTTGAAATGC ATTATTNNGA ATGTGAAGTA AATACTGTAC CTTTAAA 167

35 SEQ ID NO:7058

SEQUENCE LENGTH:69

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS08294

SEQUENCE DESCRIPTION:

GATCACATAC CCTTCTTTTA AGTGAATTTT TTTCATTTGA TTTGTCAATA AACGAATCAA 60
 ACTGGNAAA 69

45 SEQ ID NO:7059

SEQUENCE LENGTH:222

SEQUENCE TYPE:nucleic acid

50 TOPOLOGY:linear

CLONE:HUMGS08295

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EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCTCCCTG CCTGGGCTGG CTGTGAGGCC ACCTTTGTCC CAGGCCCAGC CTCAAGGCAA 60
 GGAGGGCGCT TCACTGAGGT GTGAATTGTA CGTACAGGCT TTTTATATAC CAAAAGTATT 120
 TTTTGACTAG ACCATTCAAA GCTACCCGAA CTATGTTGGA AATTTTTTTT TTCTCATTAA 180
 AATACAGGCC CTTAGGCTCT ATTTTAAATG TATNAGTCGC AN 222

SEQ ID NO:7060

SEQUENCE LENGTH:145

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08296

SEQUENCE DESCRIPTION:

GATCAAAATA CCTGTTTATA TATCTGTTAT TTAAAGAAAA ATTTTCCACA GTGTTTGTGTT 60
 ATTTTCAGC GTTCTTAGN ACATTTGGGN ATTNCACTAA AATATAGCTG GGTGGATTGA 120
 AGTAAAGGGC AANTGCGTTT GCAAA 145

SEQ ID NO:7061

SEQUENCE LENGTH:20

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08297

SEQUENCE DESCRIPTION:

GATCTTCCAA AACACACAAA 20

SEQ ID NO:7062

SEQUENCE LENGTH:227

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08298

SEQUENCE DESCRIPTION:

GATCAATCAA GAATAATTTT AATTTTCTTT TGTATTGAA ATGTAAATAG TTTTCTTTTC 60
 GATTAAAAAA ATTTCTATA ACTGCTAAAC AGTTAAANNC TTAAAGTAG TAAATGAGTT 120
 TATAGAAAGC ATGTATTCTT GATTTTNGTG CCTTGGTAAA GTTGATAACT ATTTATGAAT 180
 ATTTGACCAA ATTATTCCAG CATCAGAATA ATAAGCAAAA TAACCNN 227

SEQ ID NO:7063

SEQUENCE LENGTH:209

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08299

SEQUENCE DESCRIPTION:

GATCACACCA CTGCATGCCA GCCTGGGCGA CAGAGTGAGA CCCTGTCTCA GAAAAGAAAA 60
 AAAAAGTAAA AATTGCATGT AAGTTGACCC GCACTATTCA AATTTGTGGT GTTCAAGGTT 120
 CAACTGTAAT TTCCTAGCAG CATTTTGTGT GTTTGAGAAT CTCTTGACAC TCTTCAAGTA 180
 AATCCCTAAA TTACAACCTT GACATCAAA 209

SEQ ID NO:7064
 SEQUENCE LENGTH:130
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08300
 SEQUENCE DESCRIPTION:
 GATCGACTGA GCTCAAATTT CAGTTGGGGC CCTATTAGTC TGCCTTTCCA AAAAGGGCGT 60
 10 TGGGACTTTA ATTTAAAACT TAACCAGTTG AAAAATAAAG GTGGCGTGGA AATGCTTTTT 120
 GATTTGCAAA 130

SEQ ID NO:7065
 SEQUENCE LENGTH:223
 15 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08301
 SEQUENCE DESCRIPTION:
 20 GATCTGGGCT TTAAAGCAGA GGCCATGTCC TTGTCTGGTC CTGCTTNTGG CTACAGCCAC 60
 CCTGGAACGG AGAAGGCAGC TGACGGGGAT TGCCTTCCTC AGCCGCAGCA GCACCTGGGG 120
 CTCCAGCTGC TGGAACTCTA CCATCCCAGG AGGCAGGNAC AGCCAGGNNG AGGGGAGGAG 180
 TGGGCAGTGA AGATGAAGCC CCATGCTCAG TCCCCTCCNA TNN 223

25 SEQ ID NO:7066
 SEQUENCE LENGTH:260
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08302
 30 SEQUENCE DESCRIPTION:
 GATCGGAGAC ATCCTGCTGT TCGGNACGTT GCTGATGAAT GCCGNGGCGG TGCTGAACTT 60
 TAAGCTGAAA AAGAAGGACA CGCAGGGCTT TGGGGAGGAG TCCAGGGAGC CCAGCACAGG 120
 TGACAACATC CGGGAATTCT TGCTGAGCCT CAGATACTTT CGAATCTTCA TCGCCCTGTG 180
 35 GNACATCTTC ATGATGTTCT NCATGATTGT NCTGTTCGGC TCTTGAATCC CAGCGATGAA 240
 ACCAGGAACT CACTTTCCAN 260

40 SEQ ID NO:7067
 SEQUENCE LENGTH:154
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08303
 SEQUENCE DESCRIPTION:
 45 GATCGAAAGA AGNACATCAT GAAAAATAAA TGAACCTTGC TTAGTGGATT GACTCCTTTG 60
 CTGAAGTCAG TTATTCATCA AGAATGCAAT TAGNCTAATT GTGAATAAAT GATTGNNTGA 120
 AGNTATAATA AATAAAACCG TAAACTCTGG CAAA 154

50 SEQ ID NO:7068

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EP 0 679 716 A1

SEQUENCE LENGTH:29
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
5 CLONE:HUMGS08304
SEQUENCE DESCRIPTION:
GATCAATAAA ATGTGATTTT NCTGATAAA 29

10 SEQ ID NO:7069
SEQUENCE LENGTH:74
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08305
15 SEQUENCE DESCRIPTION:
GATCGTCCCA TCACAGTGTT GTTAATGTTT GCTGTATTTA TTAATTNCT TAAAGTGAAA 60
TCTGAAAAAT GAAA 74

20 SEQ ID NO:7070
SEQUENCE LENGTH:264
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08306
25 SEQUENCE DESCRIPTION:
GATCTGGCTT CACAGACAGT TGACACATCT AAAAGAATCT TTGTTTCCAA CTGGACTGAC 60
TCATTGAGAG GAAACCTTCA CTGCACCTCA CCAGTGCAGA CAGAAGACCT GGAGGCAGGA 120
CGGGAGGCAT TTTATACTTC AGCACGATAT GGGCCCTGAC TTTTCATTTT CTACATGGAG 180
ACATATTGCT GGAGGCTCAT CATCCTGAGG GGAGAAACAT GGATTTCATG GTTTTGCCAT 240
30 AAAAGTATGC AACAAAGAGT TCCN 264

35 SEQ ID NO:7071
SEQUENCE LENGTH:27
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08307
SEQUENCE DESCRIPTION:
40 GATCANTAAA AAAAATAAAT AATTAAA 27

45 SEQ ID NO:7072
SEQUENCE LENGTH:175
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08308
SEQUENCE DESCRIPTION:
GATCTNTTTT NTCTGGAGTC CTGGAGTTNC TTTCTAAGG TTTAATATAA GCAAGCNCCA 60
50 GAAAGAATGC TGACAGAAAA GGGACCCTAG CTGTGGTAGG AAGTGGCCCT CAGAGTCAAG 120
GAGGCAGGAT GANTTTAAAT NCTNCATGTA GGGCATATTT TGGGGAGTNA TGGGN 175

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SEQ ID NO:7073
SEQUENCE LENGTH:281
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08309
SEQUENCE DESCRIPTION:
GATCGACTTT AAGTTGCTTC GGGTCTAGTA GAAATCCGGT TATTTCCAAG CCTGTNAAAT 60
GTGAAGTTCA AGAAAATTGA ATCCGTGTAT AGCAAAACCA TTTTNTCTCA CAATTTITTA 120
ACCAATGAAC ATTGTGTGGG AAATTNATTC CATATGTTTT CAAGTCATTT ATGCAACTTG 180
AATAATTAAA ATATTCATTG GTAAGAGCTG CCTGATGTCT TTAAGTCAG GAAATTGTCT 240
TATGCCAAAA GTAGACAANC TCAAACACCA AAAGGTCANN N 281

SEQ ID NO:7074
SEQUENCE LENGTH:106
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08310
SEQUENCE DESCRIPTION:
GATCATTTTT CTTGGAGGNT GGAGATTGGC TAGTACCTCT GGCCTAACTG TGTAGGTCAA 60
TACTCTTTTA CATTGCCTTC TAATAAAAGC AGNATGATNC AGCAAA 106

SEQ ID NO:7075
SEQUENCE LENGTH:69
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08311
SEQUENCE DESCRIPTION:
GATCTTGTGT CTGTTGTGCC TGAATTTGGC GTGTCTAATA AAGGCCCATG CACACTATAG 60
GCACTCAAA 69

SEQ ID NO:7076
SEQUENCE LENGTH:292
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08312
SEQUENCE DESCRIPTION:
GATCAGTGAA ATNTTAACT CCCACGGTAA CATCATGCCT ATGGACATTA AAGGAATAGC 60
AAGGACTTGC TCCCACGACT TCACATTGC TCCCACTGGC CAGGAATGTG GGCTCTTAGT 120
GATTCCAGAT AAGGGCAAAG CTGGGGCCCA ATGTATGTAG CCAAATTGNA GNAATCATGG 180
NGCAGAGGCA AATGGTGACA TGAATTTGTA ANCTGTGTTT ATGTGTTNTT ATATTTATAT 240
TTCTGANCTC AGTACATGTT AATATTTAAN TAATTATGCA GTAACTTTCA AA 292

SEQ ID NO:7077
SEQUENCE LENGTH:29

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08313
 SEQUENCE DESCRIPTION:
 GATCCTGCAC TCTAGCACAT GACTCCAAA 29

SEQ ID NO:7078
 SEQUENCE LENGTH:334
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08314
 SEQUENCE DESCRIPTION:
 GATCCCTGT CCCCTGGGGC TGGTGGATAG GGCAGAATCC TGGGCCCCCA GAGACCTTTG 60
 CCCACACACA CTCCTTCCCC TTGTCCCTGG GGNACTCCCC CAGGATTGTG CAATAGTCAG 120
 AGTGTCCCTT TTTGCAGGGG ACTGGGCCAT GGGTCCTAGG CCCATCTGTC CATCCTCCTC 180
 TCCATGCAAG TGCTGTTTGG GCAGNAGTCA CCATGCAAGG GTGNCATCGN CANCCACGTA 240
 CCAAGNCACC GCAGCTGCTG CCACTCTGCT GCCTGTACAG AAGGAANCTG NATCTTTTNN 300
 ATATTCTAAT AAAATNACTG TGGGNTTNA NAAA 334

SEQ ID NO:7079
 SEQUENCE LENGTH:245
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08315
 SEQUENCE DESCRIPTION:
 GATCCCGTAA ATTGAAGGCA ACAAGGTCTT GGTCTNCAG TAAGGGGGGC ATGGAGCCCT 60
 GGTGGGAAGG AAAGAAGNGA AATTGTGTT CCTGATTCCT TAATCCCAN GAAAGGTCCC 120
 TCTGCTAACA TTCACATATC TNTCCTAGC CAAAGTCTGT CCATTNAAAC CCTNTAGCAT 180
 CTGAGCTATC TGCATCCAGN TTTCAAANAT ATTCGNTGT TACTAACAT GTNTGTNAGC 240
 CTNTN 245

SEQ ID NO:7080
 SEQUENCE LENGTH:88
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08316
 SEQUENCE DESCRIPTION:
 GATCACAAAT NACTTGTGA AATGTAAGGC AGTCCCCCTC CTCCTCTTTA TCTACATTAC 60
 TTCCCGAAAA TAAATGCANA TTAATAAA 88

SEQ ID NO:7081
 SEQUENCE LENGTH:149
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08317

SEQUENCE DESCRIPTION:

GATCAGGTCA CAGTGTGTAC GTAAGAGTCA GCTNCTTATG GNTATTGTCA TCTAGGTAAC 60
 TTTGTAAAA AACAAACAAA ACTGGCATTG TTAATTCCAA AAAGAAAAAA AAAAGCAGTA 120
 AAAANAAANG GNCATTTGNA CCNCCGGCN 149

SEQ ID NO:7082

SEQUENCE LENGTH:312

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08318

SEQUENCE DESCRIPTION:

GATCTCAGCC TCGGGTGGTC AGGGTGGGAA ATGATAGGAA GAAGCTGTCA TCTGCATCCT 60
 AGTTTGCCCTG AAATGAACCC AAATAATACC CATTATTATT AGTCCTGAAT TATGAGTAGT 120
 GAATGATACC CATCATTCTG GCATCATGAT GAGTAGTGTG CACTTCCATT CTGAAAAGTG 180
 CCCTGCTGTG AAAAATAAAT TATATAGTCC TCCTAGGTAA ATGAAGGAGG AGGGAGAAGT 240
 GTGAAAGAGT ATGGCTTAAA TCAGACAAGA TATACAAGAA GATACTTTAT ATAGGGCAGG 300
 AGCGGTGGCT TN 312

SEQ ID NO:7083

SEQUENCE LENGTH:87

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08319

SEQUENCE DESCRIPTION:

GATCTGGGAA ACTGATTGAA TAATACACTT TTCTTGCTTT GGTGCTCAAA GTGGTTTTTT 60
 TCCCCAATA AAATTATTTA ATTGAAA 87

SEQ ID NO:7084

SEQUENCE LENGTH:312

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08320

SEQUENCE DESCRIPTION:

GATCAGGAAA TGCATCTNCG CAGATGGTAT TTCCTTCAGA AAAGACTTTN CTACTTTAAA 60
 TATAAATNAN GCCATAACAG TTTCATGCTG TGGAAAGAGG GTGAAAAGGT TCATTTTAAAG 120
 AGATTATATA ATANGACCTT NCACATTTC TGTGAAATGT CTAACTTTNC CAGTNCTTCA 180
 GCAAGTTTTT TTGGGGGGTG ATGGGGAGNN GGTAGTATTG GTTTTAGAGG TTCAAATCT 240
 GTGANCTTTG GAGAGGGGAC AGTTGTTGGC TCTGGTATTT CCTAGTTTTG TAGTAACGTT 300
 TTGCTAGCCN NN 312

SEQ ID NO:7085

SEQUENCE LENGTH:168

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08321

EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCTGATGG TTTTAAAAAG GGGTGTCTCC CTGCACAAAC TCATTCTCTG TTTGCCTACC 60
 GCCATCCGCT TAAGATGTGA CCTGCTCCTC CTTGCCTTCC GCCGCGATTG TNAGGECTCC 120
 5 TCAGCTATGT GGAAGTGTAA ATCGAATTAA ACCTCTTTCT TTTGTAAA 168

SEQ ID NO:7086

SEQUENCE LENGTH:146

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08322

SEQUENCE DESCRIPTION:

GATCCATGTG TTTTACTCTA AAGTGCTAAA TATGGGAGTT TCCTTTTTTT ACTCTTTGTC 60
 15 ACTGATGACA CAACAGAAAA GAAACTGTAG ACCTTGGGAC AATCAACATT TAAATAAACT 120
 TTATAATTAT TTTTCAAAC TTAA 146

SEQ ID NO:7087

SEQUENCE LENGTH:314

20 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08323

SEQUENCE DESCRIPTION:

GATCAATAAA GCTGTCCTAC ACTTGTGCTA GGNAGGCACC GCTGTGACTG GGCTCCGGGC 60
 25 CTTTCCACTA CGTTGAAGAA GAAAACCTAT TTTTAAATGT AAATAAAATA TCTGGTAGCC 120
 TGTGTGGAAG GCTGACCGTT TTAAGAAAGT GCATGTGCCT TGAAAGGGGG CAGAATGTTT 180
 AGTCGGTCGT GTTTTAAACA CAGAGTCTCT AGAAGAGGTG CAGACATCCC GTCTGACTGT 240
 CCCTGTGGAC TCTCTCAGTT GTATGTTGCT ATAATCCTCC AAATCAAAGC TCTTTCTGCT 300
 30 TGTGCAAGAT TNNN 314

SEQ ID NO:7088

SEQUENCE LENGTH:71

35 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08324

SEQUENCE DESCRIPTION:

GATCTTACTG AATTTATTTT GTAAATGCCT AATGAGGCAG ATTTTGAAT TAAAGAAATG 60
 40 CTACATGTAA A 71

SEQ ID NO:7089

SEQUENCE LENGTH:412

45 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08325

SEQUENCE DESCRIPTION:

GATCCTAACA TATCAGGACA TGGACTCAGG ACCTGCCCCG TGATGCTGTT GATTTCTCAA 60
 50 AGGTCTTCCA AAACCAACA GAGCCAGAAG TAGCCGCCCG CTCAGCGGCT CAGGTGCCAG 120

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EP 0 679 716 A1

5 CTCTGTTCTG ATTCACCAGG GGTCCGTCAG TAGTCATTGC CACCCGCGGG GCACCTCCCT 180
GGCCACACGC CTGTTCCAG CAAGTGCTGA AACTCACTAG ACCGTCTGCC TGTTCGAAA 240
TGGGGAAAGC GTGCGTGCGC GTTATTTATT TAAGTGCGCC TGTGTGCGCG GGTGTGGGAG 300
CACACTTTGC AAAGCCACAG CGTTTCTGGT TTTGGGTGTA CAGTCTTTGT GTGCCTGGCG 360
AGAAGAATAT TTTCTATTN TTTTAAGTCA TTTCATGGTT TCCTGGTCCT GN 412

10 SEQ ID NO:7090
SEQUENCE LENGTH:91
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08326
15 SEQUENCE DESCRIPTION:
GATCTATTTT TGGCTTATGT CTTTAAATAC TACTCTCTTT CTCTGAATTT TGAAATATGA 60
AGTAAATCT AGCCCATTTG TTTTATGCAA A 91

20 SEQ ID NO:7091
SEQUENCE LENGTH:294
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08327
25 SEQUENCE DESCRIPTION:
GATCTCAAGA GTTCAAGACC AGCCTGGGCA ATATGGTGAA ACCCGTTTGT ATTAAAAATA 60
CAAAAATTAG CTGTGCATGG TGGCCACGG CTGTAATCCC AGCTACTCAG GAAGCTGAGG 120
TCCGAGAATC ACTTGAATGC TTGAACTCGG GAGGCAGAAG TTGCTGTGAG CCGAGATGGC 180
ATCACTGCGC TCCAGCCTGG GCAACAGAGA GAGACTGTCT CCACTGTCTC CAAAAAATA 240
30 NGAAAAAAGG AAGTTGTTN GATTCTNTGGC CCTCAAGGAA ATTNTGGAAA ANAN 294

35 SEQ ID NO:7092
SEQUENCE LENGTH:79
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08328
SEQUENCE DESCRIPTION:
GATCATGTTT TGAAGCAGCA GGTCCAGGTC ACTTTGTATA TAGAATTTTG CTGTATTCAA 60
40 TAAATCTGTT TGGAGGAAA 79

45 SEQ ID NO:7093
SEQUENCE LENGTH:122
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08329
SEQUENCE DESCRIPTION:
GATCTTTCCC TAGAGATGT AACATACGTT TTTATTCATT TAATCACTTC ATTATGCCGG 60
50 GGTTAATNAT GTTATTTTA TAATTGGTAA TAAAGGCCAC ATTTATTTT GTAACGTGTA 120
AA 122

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SEQ ID NO:7094
 SEQUENCE LENGTH:50
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08330
 SEQUENCE DESCRIPTION:
 GATCTTAAAC TCTTTGATTA AAAATAATAA ATGGAAATGT GTTAAGCAAA 50

SEQ ID NO:7095
 SEQUENCE LENGTH:77
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08331
 SEQUENCE DESCRIPTION:
 GATCCAGGGA TTTGGGGAAA AGGAAGGAAT TTGATGTTTT TTGGGGTGGG AGGGGAGGGT 60
 GTGTTTTTTA CACCAAA 77

SEQ ID NO:7096
 SEQUENCE LENGTH:186
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08332
 SEQUENCE DESCRIPTION:
 GATCGCTGGC CTTCTGACT TAACATTNAT GCATGCCAAG GTGTTGATGT TCCCAGCCAC 60
 GTTAACACCT TCCACAAGNT CTCAAGAAAA AGCAGACGGA TAACTGATGT GAATTGGACA 120
 GTTTCTATTG CTTTTCTTT TTTCCATCCC TTCCCTACCA TCAAAAGCAT ACCTGCTCTA 180
 ATTAAA 186

SEQ ID NO:7097
 SEQUENCE LENGTH:291
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08333
 SEQUENCE DESCRIPTION:
 GATCATATAC TCTGCTGTGG TCATGCCCAC TCTTTGGGAG TATATCCCT TTATATATAT 60
 TGAGTATTGT ACCACTTGAG AAATTCCTTT GTTCTGTTAT ACAAATTAA TCTTTCTGCT 120
 CATAATGATT GATGATACCA CCAGTAAAA TAGGATGTTT ACCCCAAAAC AAGTGCAAT 180
 TAAGAATTG AACACAACCA CATTTTTTAA AATGAACTT CTATCGGAAG TAAATTAATT 240
 TGNGGTAATA AAGTCCAGTA TTNGGNAAA TGTACAATGT TAAATCTCAA A 291

SEQ ID NO:7098
 SEQUENCE LENGTH:307
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS08334

SEQUENCE DESCRIPTION:

5 GATCAGACTT TTGTNTCTTG GGTCCCAGAT GGCACAGGAG CACTGCATGC TTGTTTTCTA 60
 GAGCCCAGCC AGTCATGGGT GCTAGCCTAG TCTCCACACA CCAGCAAGTA GAACCCAAGT 120
 GTANNNNNAT AAATATTTC TGAGTACCAG TAAGAGAATG CATTCTTTTC TCATCTAGGC 180
 CAGGAATGTT GAAAATGCTC AGCCTTACAT AGAACTCCT AGATTTTCAC TAACGCATTT 240
 CACAAAAGTA AATAAGTATT TCATATAATT CAGAGGATGT TTAAATTGTC AGCATTTTAA 300
 TAAATAN 307

SEQ ID NO:7099

SEQUENCE LENGTH:199

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08335

SEQUENCE DESCRIPTION:

20 GATCTGTACT TTGAACAAAG AAAGTGTGTA GAGACCCCTG GGAGTGAGGG GTAGGGGATA 60
 CAATACTCTG AAGCAGAAGA GGACCCCTGC AGTGATGGTT AGGGACTCTT CTGAGTAGGA 120
 GACGTAATCG GCTTTTGGTC CATGGAACCT TTACCACATG TATTCAGCAT ATATACACGA 180
 TTAAAGTGCT GCAAATAAA 199

SEQ ID NO:7100

25 SEQUENCE LENGTH:297

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08336

SEQUENCE DESCRIPTION:

30 GATCTGGGAG TGGCTTCCTA ACGGAAATGG AACAGGATTA CCAACTAGTN GACGAAAGTA 60
 ATGCTTTCCA TGACAACCTT AGGTCTCTTG ACAGNAATCT GCCCTCAGAC AGCCAGGACT 120
 TGGGTCAACA TGGATTAGAA GAGGATTNA TGTATAAAA GAGGATTTTC CCACCTTGAC 180
 ACCAGGCAAT GTAGTTAGCA TATTANNGT ACCATGGTTA TATGATTAAT CTTGGGACAA 240
 AGAATTTNAT AGAAATTTT AAACATCTGA AAAAGAAGCT TAAGTTTAT CATCCTN 297

SEQ ID NO:7101

SEQUENCE LENGTH:296

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS08337

SEQUENCE DESCRIPTION:

45 GATCTTTGTC TGTGAAGAAG AAAATTATCT CCCTAGTTCA ATCTGTAGTG AAATAAGGCT 60
 ACAGAAGGCA TTGTTTTTTC CTTTTNATT TTTTGTATTA TATATTNCC TTAAATATGT 120
 TTTATTGTCT TCTCTANGCA AAAAGTTCTT AATAANCATA GTATTTCTCT CTGCGTCCTA 180
 TTTCAATTAGT GAAGACATAG TTCACCTAAA ATGGCATCCT GCTCTGAATC TAGNCTTTT 240
 AGAAATGGCA TATGGTTTTG ATGATATGNC CACCATCCAA AATGTCCGAN TNAATN 296

50 SEQ ID NO:7102

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SEQUENCE LENGTH:305

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08338

SEQUENCE DESCRIPTION:

GATCAACTTT AAGAAGATTT AAAAAACGCT CAAGAAAGAN GAGGATATCC NACTNTTCCC 60
CGTGCAGACA AAATACATGG ATGTGGTCAA NGAGTGAATC CGTTTAGCTC GACAGATTGA 120
GAAATCTGAG TATCGGAACT TCCAGGCTTG CCTNCACAAC TCTTGGATTG AGCAGGCAGC 180
AGCTGCCCTG GAGATTGAGC TGGGAAGANGA CATGTATAAG GNAGGAAAAG CTNNCCAGCA 240
AGAAGAACGT CGGNGACAAA AGCAGATGGN GGTCTNGAA GAAGGAGCTG CGNCNCCTGC 300
TTNTN 305

SEQ ID NO:7103

SEQUENCE LENGTH:414

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08339

SEQUENCE DESCRIPTION:

GATCCACAAG TCCTTGTTC ACTGTGCCTT GGTTCCTCTT TTATTTCTAA GTGGAAAAAG 60
TATTAGCCAC CATCTNACCT CACAGTGATG TTGTGAGGAC ATGTGGAAGC ACTTTAAGTT 120
TTTTCATCAT AACATAAATN ATTTCAAGT GTAACCTATT AACCTATTTA TTATTTATGT 180
ATTTATTTAA GCATCAAATA TTTGTGCAAG AATTTGAAA AATAGANGAT GAATCATTGA 240
TTGANTAGTT ATAAAGATGT TATAGTAAAT TNATTTTATT TTAGATATTA AATGATGTTT 300
TATTAGGATA AATTTCAATC AGGGTTTTTN GGATTAANCA ACCANCCAAT TGGGTCCCA 360
GTAAANTTTC CATTTTCAGAT ANCCACCANN TAGTTTCNNN GTNTAGGTCC ATTN 414

SEQ ID NO:7104

SEQUENCE LENGTH:252

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08340

SEQUENCE DESCRIPTION:

GATCCCCTGA CTGGCTGGG GCAGCTCCCA GGATATCCTG CCTTCCAAC TTTCTGAAG 60
CGCCTCCTCC TAACATGGCG ATCCGGAGG TCAAGNNCNT GGGCTCTCCC CAGGGTCTAA 120
CGGTTAAGGT GACCACATAC CAGTGCCAAG GGGNTNTAA AGTGGTGATG TAGTTNTGCT 180
CCCCTCCCA AGAGCGGGTG GNGGGGGTT NAANATGTTT GGCCTGNNTA AGTGGGCCTT 240
CCCATTNAAG TN 252

SEQ ID NO:7105

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08341

SEQUENCE DESCRIPTION:

GATCTTATTT TTAAATTTTA AGTGCCACTA TTAATGTAAA AAGGGGGGGG CTCTACAGCA 60

EP 0 679 716 A1

5 GTCGTGATGA AACTTAAATA TATATCCTTT GTCCTCGAGA TTTTAGGAAG GGTGTAGGGT 120
 GAGTAGGCCA TTTTNAATTN CTGAAGTGCT AAGTGTTTT ATACAGCAAA CAAAAAGTCA 180
 ATTTTCCTTT CCACCAGTGC GAGAGAGGAT GTATACTTN CAAGAGAGAT GATTGCCTAT 240
 TTNCCGTTTG ACAGAGTCCC GTAGATGAGC AATGGGGAAC TGGTTGCCAG GGTCTAANTT 300
 TGGATTGATT TATGCACN 318

10 SEQ ID NO:7106
 SEQUENCE LENGTH:299
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08342
 SEQUENCE DESCRIPTION:
 15 GATCTGTGAC CCCAGCCATG AGGACCCTCG CCATCCTTGC TGCCATTCTC CTGGTGGCCC 60
 TGCAGGCCCA GGCTGAGCCA CTCAGGCAA GAGCTGATGA GGTGCTGCA GCCCCGGAGC 120
 AGATTGCAGC GGACATCCA GAAGTGGTTG TTTCCCTTGC ATGGGACGAA AGCTTGGCTC 180
 CAAAGCATCC AGGCTCAAGG AAAACATGG CCTGCTATTG CAGAATACCA GCGTGCAATG 240
 20 CAGGAGAACG TCGCTATGGA ACCTGCATCT ACCAGGGAAG ACTCTGGGCA TTCTGCTTN 299

25 SEQ ID NO:7107
 SEQUENCE LENGTH:302
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08343
 SEQUENCE DESCRIPTION:
 30 GATCTTAGAG GTTGAAGCCC ACCTTCTCTT TTCACATAGG AGGGAACAGA CCATGGAAAT 60
 TTAAACGACT TCCTCACGGT CACAGAACTA GTTTTTTAAT CCTCAGGCAG TGNATCCCCC 120
 CACCCTACAA CTGTGCACAA CCTCTTTCCC CACAGTGCAA TTCAGAATAT GCTCAGGGAA 180
 TGCCAGGCAC CTTGTAAAC TGCTGGGAGA AAAGCATGNT TCCCACAAGG ACTAAGTATC 240
 AGTGATTGT AATTTCTCTG TTTTGATTA TCTGCTTTGC TGATGTAGAC AAGAGTTAAC 300
 TN 302

35 SEQ ID NO:7108
 SEQUENCE LENGTH:110
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 40 CLONE:HUMGS08344
 SEQUENCE DESCRIPTION:
 GATCNACTG TCTCACTGCT TTTCCAATA TAATCATNTC GCTCTNACTT TAACAAAAGG 60
 CGATGGCACA GTAATTTAGT AAATTTNTGT AATAACATNT AAAAAAAGN 110

45 SEQ ID NO:7109
 SEQUENCE LENGTH:121
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 50 CLONE:HUMGS08345

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EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCCTGTCT AAAAAAGAAA AGAAAAGAAA AGAAAAGAAA AGAAAATCAG CATTGTATCC 60
 CCAGAACCTG CTACAGTGTA CTGTACACTT AATAAATATT GGTGAATGA ATGAACTGAA 120
 A 121

SEQ ID NO:7110

SEQUENCE LENGTH:99

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08346

SEQUENCE DESCRIPTION:

GATCATCAGT AAATTTGTCA TGTTATATAT TTATTTTTTT ATAAATCAAG ACTTCTGTGT 60
 GCTCTTAAAT ATATTAAAA CAATTTACAT TTCAGGAAA 99

SEQ ID NO:7111

SEQUENCE LENGTH:92

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08347

SEQUENCE DESCRIPTION:

GATCCAGNA GGAGGGAAGA AAATTTGCAG TGAAGAAAA CAGTAAAAAA AAAAAAATA 60
 ATGTATAAAA AAGTTGCATT ACACAGTNCA AA 92

SEQ ID NO:7112

SEQUENCE LENGTH:304

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08348

SEQUENCE DESCRIPTION:

GATCATTCTN GTTGATACCC TTCTTTGAAT ATTCTAGTGT ATTAATATAC CATTGGTTAT 60
 TNAATCATGT CTNATTAATG GACTGGCTGT TTTCACATAT TNGATATATC AAGTGTCTTC 120
 ACAACTTGTG CTTGCATATT CTTTCCCAA ATATTGAAAG TCCATATATT TCCTTGATCA 180
 TNTTTAAAGT TGATATCTNA ATNTNTCATT GTAGTNGCAA AGCATGTAAT TNCTTGGGGG 240
 AGGGGGGCTG TAAATATTGA CATTTTAAAA TAAAACTTTT AAATCAGCCT TAAATGTATC 300
 AGGN 304

SEQ ID NO:7113

SEQUENCE LENGTH:309

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08349

SEQUENCE DESCRIPTION:

GATCAAATTT CACGTCGTCT TCTGTATACT GTGGAGGTAC ACTCTTNTAG AAAGTNCAAA 60
 AAGTCTACGC TCTNCTTTCT TTCTAACTCC AGTGAAGTAA TGGGGTCCTG CTCAAGTTGA 120
 AAGAGTCCTA TTTGCACTGT AGCCTCGCCG TCTGTGAATT GGACCATCCT ATTAACTGG 180

EP 0 679 716 A1

CTTCAGCCTC CCNACCTTCT TCAGCCACCT CTCTTTTCA GTTGGCTGAC TTCCACACCT 240
AGCATCTCAT GAGTGCCAAG CAAAAGGAGA GAAGAGAGAA ATAGCCTGCG CTGTTTTTTT 300
AGTTTGNNN 309

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SEQ ID NO:7114
SEQUENCE LENGTH:282
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08350
SEQUENCE DESCRIPTION:
GATCTTCCCC AGCTATGAAG CCTATCTTGG AATGGGAAAC CATGAAATTT GTCTTTTAAA 60
AGAAAGGGAA GAAATAGCCT TACAGCCCCT GTTCCTCCAA GTGCACATTT AGCCTTAAAA 120
TCCTGGGAGT CGGGAGACAC CGTGGAGTCC TTACATCCAC AGTGTCTGCT GGTCTGGGTC 180
AGAAGGCAGT AATGCAAGAG TCCTTTTGTG AAGAGTGTTT CTATGTAGAG ATGTTTATAT 240
TTAAGTAGTT CTTTATAAA TAAAAGCATT TCTAATGGCA AA 282

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SEQ ID NO:7115
SEQUENCE LENGTH:27
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08351
SEQUENCE DESCRIPTION:
GATCTGTTTT CTGTAACTTT TGCTAAA 27

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SEQ ID NO:7116
SEQUENCE LENGTH:251
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08352
SEQUENCE DESCRIPTION:
GATCTTAGGC TTCAGCTCTT TATCTTACAC CTTGAGACCC CAAGAGAGTC CCAAATTCCT 60
GGAAAGGCAG CATCACAGAG TGGGAAATA CAGGACTTGG CTTAGGTTC TAGTCCTTAA 120
CACAGCTTCT TCCTTGCTGT GTGTTCTTG GTAGTCACT AAATCTCTCT GAGCCTCAAA 180
TTTCTTATCT CTAGGTTGAA AAATAAAATA AAGTCCCCT GCCAATCAGT GCAAGGAATC 240
TGTGAGATAA A 251

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SEQ ID NO:7117
SEQUENCE LENGTH:123
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08353
SEQUENCE DESCRIPTION:
GATCACGCCA CGGCACTCCA CCCTGGGCGA CAGAGCGAGA CCTTGTCTCA GAAAAATCT 60
GTGAGGTAA AAGGCTGTGT GTGTGAAAT ATCTTGAAA TTGTAAAGAA CTACACAANT 120
AAA 123

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SEQ ID NO:7118

SEQUENCE LENGTH:295

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08354

SEQUENCE DESCRIPTION:

GATCATTCAT CAACATGTGT CTCTNAATTT AATTCAGTAG TTGTAATTAT TCTTTGGTTT 60
 AGACCAAGAA AAAGGAAATC CCCCTTTNA ATGTATTCCT TGGTTTNAGG ACATGACTCC 120
 TGTAAAGNAG AGGAAAGGGA GATGCTTCCT GTTTGANCTG CAGTGAATTC ACGGTTCCCTG 180
 TTTCACCACT CCAAACCTTA TGGCGACTCA CACACACATT CCTCTTNTCT GTTACTGCCA 240
 AAGGTTCCGGG TTTAGTACAC TTCAGTTCCA CTCAAGCATT GAAAAGGTTT TCGTN 295

SEQ ID NO:7119

SEQUENCE LENGTH:203

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08355

SEQUENCE DESCRIPTION:

GATCCCCAGG ATGCCCACTT CCCTCCACCC CCACACCCCA GCCGTGTCCC TAACCCCTGG 60
 CAACCAGGAA TCCACTCTCC ATTTCTATAA TGTGNNATT TCAAGAATGT TATTCAATGG 120
 AATCATATAG TATGTAACTT GTTTGAGCT TAAAAAATAA GTATACATGA CTTTAATGAG 180
 GAAANTAAAA NTGGATATTG AAA 203

SEQ ID NO:7120

SEQUENCE LENGTH:82

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08356

SEQUENCE DESCRIPTION:

GATCCGTTAG TGTCTGTTTA GAAAAGATGT TATAAACTTA CAGAAACAAA TATAATAAAC 60
 TGAAGCAGAT TTGAAAAGCA AA 82

SEQ ID NO:7121

SEQUENCE LENGTH:79

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08357

SEQUENCE DESCRIPTION:

GATCTTTTGC ATTGCAGAGA TTTTATATAT ATTTTGTTTG TAATGAGCCA TTCTCAATAA 60
 ACATTATTCT CACTGCAAA 79

SEQ ID NO:7122

SEQUENCE LENGTH:299

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08358

SEQUENCE DESCRIPTION:

5 GATCCCTAGT AAACCTCTTC TTCACCTTTA CTGTCAGATT TACAAAGGTC CTCCCATTGC 60
 AAAGCAGTGN NNNTCCTAAT TTATATATTG TTTTNCCTAGT TCATTTTGTG TTTCCAACTT 120
 TTCATGTAAA ATTTTAATTA TTTTGAATG TGTGGATGTG AGACTGAGGT GCCTTTTGGT 180
 ACTGAAATNC TTTTCCATG TACCTGAAGT GTTACTTTTG TGATATAGGA AATCCTTGTA 240
 10 TATATACTTN ATTGGTCCCT AGGCTTCCTA TTTNGTNACC TTGCTTNNN NGTGGCANN 299

SEQ ID NO:7123

SEQUENCE LENGTH:108

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08359

SEQUENCE DESCRIPTION:

20 GATCATGTGC CAAGGCTGCG GTGCGTTCTG TCACGATGAC TGTATTGGAC CCTCAAAGCT 60
 CTGTGTATTG TNCCTTGTGG TGAGATAATA AATTATGGCC ATGGGAAA 108

SEQ ID NO:7124

SEQUENCE LENGTH:113

SEQUENCE TYPE:nucleic acid

25 TOPOLOGY:linear

CLONE:HUMGS08360

SEQUENCE DESCRIPTION:

30 GATCAGACTG TGCTTTCTAA TTCTAAAAA TTGAGCAGTG TTGGTATTCT TNCAGTATTT 60
 TTNCTCTCAC CTGGAAACT ACCTGGAAAGT TGTCAGGCAT TTNCTTACTG NNN 113

SEQ ID NO:7125

SEQUENCE LENGTH:125

SEQUENCE TYPE:nucleic acid

35 TOPOLOGY:linear

CLONE:HUMGS08361

SEQUENCE DESCRIPTION:

40 GATCTCTAAG TAATTNTNCA AAATCCGGTA TTACTATGTC AAGTTATTGC TTTTGGTAAA 60
 TTCACTGTAC CCAGTTATAA ATGAAAGANT ATGGANTTAA AAATTTTAAA NCTAAATAAT 120
 TTGNN 125

SEQ ID NO:7126

SEQUENCE LENGTH:122

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS08362

SEQUENCE DESCRIPTION:

50 GATCATGCTC CATCCAGCCC CACCCAATGG CCTTTTGTGC TTGTTTCCTA TAACTTCAGT 60
 ATTGTAAACT AGTTTTTGGT TTGCAGTTTT TGTGTGTGTT TATAGACACT CTTGGGTGTA 120

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5 SEQ ID NO:7127
 SEQUENCE LENGTH:256
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08363
 10 SEQUENCE DESCRIPTION:
 GATCGGCACT CACGATTGTN ATGTTGGCTA CATCCTTGAC AAAGCCGATG GCCTTTCCAG 60
 GTTGGAACCTT GCAATAACCA TCCTGTTGAG GATGCAAAGG GCATGAAATA CAGGTTATGG 120
 ACCCGAAGTC TCAGCCTCCC CACGCAGTTC AATAACATTT ACTAAACAAA ACAAGATGTG 180
 15 CCAGGGCCTG GGGGATGGGA TAATTCAGA GAGAATTAAA GCATCCTTGT CCTCAAGGAG 240
 CTTAGGTCT GGTAAA 256

SEQ ID NO:7128
 SEQUENCE LENGTH:346
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08364
 SEQUENCE DESCRIPTION:
 GATCAACAGG GGTGAAAAGA ACCACCCTGA GGTTCCTATG CCTCTCCAN NTTAGTGGT 60
 25 AGNATTTTGT GTCTTTACTC CACCCTTCAC CCTAGTTCCA CCAAGGTTCA CACACCAGAT 120
 GTAACCTGTTT TNNAGCTGAG TTGTATGGAT TAACTTCAGT CCACTGTAAG TACACCTGGG 180
 ATGGGGTGGG GTTGGGGTTG TTAGGGGAGA AGCAGCCAGA CTTGCTTTGT GAACTGAATG 240
 TATTTTATG CAATTTTGAG TGGCCTTTCA ACCCTAAGAT GAATGGTTT GTTTTACTGG 300
 30 TTGTTGTTAT ATAGTTTGA GTATTCTGTG TTTGAAAGTT TGGGTN 346

SEQ ID NO:7129
 SEQUENCE LENGTH:42
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS08365
 SEQUENCE DESCRIPTION:
 GATCGTGTCT CTGATAAAG AATAAAGCT GTTTATTGCA AA 42

40 SEQ ID NO:7130
 SEQUENCE LENGTH:338
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS08366
 SEQUENCE DESCRIPTION:
 GATCTTAGGT CCCACTCANG GTCCTAACTG GCTATGGCCA CAGCTGGAAC TACCCGAGAA 60
 GNCTTTTGTA CATGTTTGGT AGCCGTAGCA CAAGTGATTG GAGTAGAACA TGCTACTGCT 120
 50 GTACATTTT AACTCCCCTA ATGGTGTGTC TATAATTNTN AAATCTAAGG GGGGGGGCTC 180
 AGTAAAGCCT CCTGGCACCA GGCCTTCTG CTCGACTGAA AAAAAATTCN TCTTTGAAAA 240

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TCCCCTTTTA CTCATGGCCC ACAGTAGAAT ATCCAAAACG CCTTGGCTTT CAGGCCTGGC 300
 CTTTCCTACA GGGAGCTCAG TAACCTGGAC GGCTCTAN 338

5 SEQ ID NO:7131
 SEQUENCE LENGTH:294
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08367
 10 SEQUENCE DESCRIPTION:
 GATCTTTTCA AGATGCATTC CAGATGAACT GCTAGGTGAG GGGGAAGCTT CATTITTTGTT 60
 ACCTGATAGA ATAGCTTTNC TNATGAGATA TATATAATGT GATACTATGT TTGGATATTT 120
 TTGGTCTTAA AGCAAGACTC AGTGGTGTAT CTNCATTAAA AGCTTCCTTT AAAAAAGNTA 180
 15 CAGGGNTACT AAAAAACAA GNACCCAAAC AATCAAGNTG GGCCACCTT GGNACCNTGT 240
 TTGNATNAC CTTCCATGGT TTGGTTGGCG GTATGGNAAA AGGATTGNNT GGTN 294

SEQ ID NO:7132
 SEQUENCE LENGTH:107
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08368
 SEQUENCE DESCRIPTION:
 25 GATCATGCTG TGTGGAGGAC AAATCTTTAA AATTCCAGCA TTGTATTGTC TATTGACACA 60
 CAAAGTTTGA AAATAAAGGG GCAAAAAATT TTAATGCAAA AATGAAA 107

SEQ ID NO:7133
 SEQUENCE LENGTH:328
 30 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08369
 SEQUENCE DESCRIPTION:
 35 GATCTGCGAT GGCTGACATN AATAGGTCAC CATGTTGGCA GCTGTATAAT CAGCAATGAA 60
 AGCTGAATAT TATCACCAGA TAACCAAATT ATGAAATATC TATATNANCT TCANGATTCA 120
 GGNCCANTGA ACCAGTATCT GCCCTTACGA GATTAGCTG AANGTGAANT GCAAGGTGGT 180
 GAACGGCATT CTAACAAGAG TGTGAAGAAA GAGATNGANG GCTGTCCGCT GCTGGAGGGG 240
 40 AAGAANGGTG GGGCGGCTTT TNCAAGGCTC TGTGGCTCAG TGGACCTTCA ATANATTGGA 300
 CTTCAATAA ATTAATTGNA AANCANGN 328

SEQ ID NO:7134
 SEQUENCE LENGTH:244
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08371
 SEQUENCE DESCRIPTION:
 50 GATCTGAGGG ATTTGCTCGT CATTGCTGT GGCAGAGATG ATGTTGACAC CCTGGGATGC 60
 TGTGGGAGGG ACAAAGACT CACGTATATG TTCTGTCAA TCGGAGAAGT AAATAATCCG 120

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5 TTGGGATGCC AGGATACTTT ATTCTACAGC GAAGTTAATT TTGTATCCGT CGAGGCTACA 180
GTAGAGGTAA AAATGACATT GAGTTGTACT TTGCTTTGAA TTAAAGTTGA TGGGTAGCAG 240
CAAA 244

10 SEQ ID NO:7135
SEQUENCE LENGTH:48
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08372
SEQUENCE DESCRIPTION:
GATCTTGACT CCAAAAATTA ATAAATAAAT AAAAATAAAT AAAATAAA 48

15 SEQ ID NO:7136
SEQUENCE LENGTH:48
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
20 CLONE:HUMGS08373
SEQUENCE DESCRIPTION:
GATCATGGTA TAATTAATTG AAATTTATTA AAATCTGTTT TTATTAAT 48

25 SEQ ID NO:7137
SEQUENCE LENGTH:267
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08374
30 SEQUENCE DESCRIPTION:
GATCTGTGAC TTCAAGTTTG GACCGTGGA GAACAGCACT TTCAAACCCA CAACTGAGAA 60
TNATGACACA GGGACAAGTA GCAGTGATAC ATCAGATGAC GACGATGTNT GAAGGATTTT 120
CTAACAGCTT TNGAAATCTT AGTGTGATAC ATCTCTCATA CAGTTTGGGG NGAATTGTNA 180
AAATGAAAGG ACTATANTTT ATGTAGTGGA ATACCCCCNN AGAAGAGGNN TTTTGGGG 240
35 GCTTCAACTT GNTGGAAACC NAGANTN 267

40 SEQ ID NO:7138
SEQUENCE LENGTH:314
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08375
SEQUENCE DESCRIPTION:
45 GATCATTGAA ATTCTGAGAA AACTTCTTTT AAACCTCACC TTTGTGGGGT TTTTGGAGAA 60
GGTTATCAAA AATTTTCATGG AAGGACCACA TTTTATATTT ATTGTGCTTC GAGTGACTGA 120
CCCCAGTGGT ATCCTGTGAC ATGTAACAGC CAGGAGTGTT AAGCGTTCAG TGATGTGGGG 180
TGAAAAGTTA CTACCTGTCA AGGTTTGTGT TACCCTCCTA TAAATGGTGT ACATAATGTA 240
TTGTTGGTAA TTATTTTGGT ACTTTTATGA TGTATATTTA TTAAACAGAT TTTTACAAAT 300
50 GGAANAAAGA NAAA 314

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SEQ ID NO:7139

SEQUENCE LENGTH:349

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08376

SEQUENCE DESCRIPTION:

GATCCAGAGT AAATGAGAAG TCTCTATCTG AGCTGGTCAG TTAAGTGGAGT ACATGTNACT 60
 AATCTGGGTT TAAAGTTTAC TTCATTATCT GCTAGTGTC TCCACAGCAG TTCATCCTCA 120
 TCCACACTAA GCCATCCTGT NAGCTTTTAA AGGAAGTTAA TTAAATTAAC ATTAATATAC 180
 TCTATGGGCT CCCTCTCCCA CCTGTCTGCA TAGAAAGGCA GAATTAGACA TAGCATGCTT 240
 TGGAAAAGCA AATAGGAATT GTTGGGAATG ATTTANTCTT GTTGTGTGTG TTGTTGTTGT 300
 TCACTTGTGG TTCTACATTC CTGGTGNATG NTGAATGTNG CTGTCAAGN 349

SEQ ID NO:7140

SEQUENCE LENGTH:347

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08377

SEQUENCE DESCRIPTION:

GATCTTGTGC TTCCATTAA GAAATTCTTC CATTTAAAGA AGAAAAAAAA TCTCTCTAAT 60
 TGAATATCTG AAGATATATG AAAAAGCCTA TGCTTTTAAA TTAANCTGTT AAGACAGTCC 120
 ATTGAAAGAT TGTGGAAGTN CACATCTATT TTNCACCTTA ATTTTTCCTA TTGTCCCTAC 180
 TCATGACTCT AAAAAGTGCA TGGCTTGGGG CTATACTTTG TTTTGCAGTT GTTTGGTATC 240
 GTGCCTTTCC TTATCTACAT TAGCTTAGAC TATACCTTNT TTTTNAGAAG GGGGAAGTGGN 300
 AATTAACGT GGCAAAACCT ATTTNGGCAC ACCNCCTTTG TTCAATN 347

SEQ ID NO:7141

SEQUENCE LENGTH:121

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08378

SEQUENCE DESCRIPTION:

GATCTTAATT CTTTGCAATT ATANATGAAA AATGACTGAT TTTCTNAAA ATATGTAAT 60
 TATNTAAATA TATCTGTTTG TACAGATTTT AACCATAAAA ACATTTTGG AAAACCATAA 120
 A 121

SEQ ID NO:7142

SEQUENCE LENGTH:409

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08379

SEQUENCE DESCRIPTION:

GATCTGGAAA CTATTTGGGT TTTGTTTCA ACTTTTCATT NGGATGTTTG NCNTTGACACA 60
 CACACATCCA CCGGTGGAAG AGACGCCCGG TGAACACACC TGCTGCTTT CTAAGCCAGT 120
 NAGGTTGAGG TGAGAGGTTT GCCAGAGTTT GTCTACCTCT GGGTATCCCT NNTGTCTGGG 180

EP 0 679 716 A1

5 ATAAAAGATN TCAAGCCAGT TGGCGGGATG GAATGGNTGC ACCGCAAATA ATGCATTNC 240
 TGAGTTTCT TGTTAAAAA AAATTTTTT AAGTAAGNAA AAAANAGGTA ATAACATGGC 300
 CAATTTNTT ACATAAAATG GCCTTCTGG NGTNTAAATT ATTCCTAAAA AAATCCNGTT 360
 TTATATANAC NATCCAGTTG NTGGAAAAA NTTTTCCAAA NGTTTTTGN 409

SEQ ID NO:7143
 SEQUENCE LENGTH:89
 SEQUENCE TYPE:nucleic acid
 10 TOPOLOGY:linear
 CLONE:HUMGS08380
 SEQUENCE DESCRIPTION:
 GATCTAGAAA ACTTTGGATT TTTGAAGTAA ATTTAATGT TTCATATTAA TTTCTTGAAA 60
 15 ATGTATTAAA TGTCATTGAA AGCCTTAAA 89

SEQ ID NO:7144
 SEQUENCE LENGTH:375
 SEQUENCE TYPE:nucleic acid
 20 TOPOLOGY:linear
 CLONE:HUMGS08381
 SEQUENCE DESCRIPTION:
 GATCACTGTG GACCATCTTA GCAGTTGACC TAACACATCT TCTTTTCANN ATCTAAGAAC 60
 25 TTTTGGCACT GTGACTAATG GTCCTAATAT TAAGCATGTT GTTTATATTT ATCATATATC 120
 TATGGCTACA TGGTTATATT ATGCTGTGGT TGC GTTCGGT TTTATTTACA GTTGCNANTA 180
 CAAATATTTG CTGTAACATT TGA CTCTAA GGTTTAGATG CCATTTAAGA ACTGAGATGG 240
 ATAGCTTTTA AAGCATCTTT TACTTCTTAC CATTTTTTAA AAGTATGCAG CTAAATTCGA 300
 30 AGCTTTTGGT CTATATTGTT AATTGCCATT GCTGTAAATC TAAAAATGAA TGAATAAAAA 360
 TGNTTTCATN NTAAA 375

SEQ ID NO:7145
 SEQUENCE LENGTH:513
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08382
 SEQUENCE DESCRIPTION:
 GATCTCGCTC CTTTGAGAGG TCCCATAAAA GCAAGCACCA TGGTGGCAGT CGCTCAGGAC 60
 40 ATGGCAGGCA CAGGCGCTGA CTTTCTCTTC CTTTGAGNCT GCATCAGTTC TTGGTNTTGC 120
 CTATCTACAG TGTGATGTAT GGA CTCAATC AAAACATTA AACGCAAACT GATTNGGATT 180
 TGATTTCTTG AAACCTCTA GGTCTCTAGA AACTGAGGA CAGTTTCTTT TGAAAAGAAC 240
 TATGTTAATT TTTTGCACA TAAAAATGCC CTAGCAGTAT CTAATTA AAA ACCATGGTCA 300
 45 GGTTC AATTG TACTTTATTA TAGTTGTGTA TTGTTTATTG CTATAAAGAC CTGGAGNGTG 360
 AATTCTGTAA AAATGTATCT TATTTTATA CAGTTAAAT TTGCAGACAC TGTTCTATTT 420
 AAGNGGTTAT TTTGTTTAA TGGATGGGNG ATNCCTTNTT AANCACTGGN TTGGNCTGGA 480
 TNGTGTAAGG TTTTNCAGG GNNTAGTTCC TGN 513

50 SEQ ID NO:7146

55

SEQUENCE LENGTH:180
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08383
 SEQUENCE DESCRIPTION:
 GATCTGAGAA TCTTTATTGA GAAAGAGCAC TTAAGAGAAT ATTTTAAGTA TTGCATCTGT 60
 ATAAGTAAGA AAATATTTTG TCTAAAATGC CTCAGTGTAT TTGTATTTT TTGCAAGTGA 120
 AGGTTTACAA TTTACAAAGT GTGTATTAAA AAAAACAAAA AGAACAAAA AATCTGCAAA 180

SEQ ID NO:7147
 SEQUENCE LENGTH:187
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08384
 SEQUENCE DESCRIPTION:
 GATCAAGAAA CTGGAAATG TGAAGCAATC TCTATTGAGA TAGCATCTTA GTGTTTGA 60
 GAAATCAAGA ATTTTAAAA ACAAGAATAT CAACATTTGG TTTGTGTAT AAGTGGTGT 120
 TGTATTAATA TACTTTTCA ATGAACGTGA TAAACTATGT TTTATTAAAC TACAATATAT 180
 CAGTAAA 187

SEQ ID NO:7148
 SEQUENCE LENGTH:133
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08385
 SEQUENCE DESCRIPTION:
 GATCCAGAGG CGAGAAGCCA CCCTGTTAGA NCTCCTGTCC ATCATCCAGG TAAGTCAACT 60
 CTCCTCTTC AGGGCCAATC CCCAGACCTT TTGTTGAGCC AGGNCNNTAT CACCTNTCCT 120
 ACTNACTTAA AGN 133

SEQ ID NO:7149
 SEQUENCE LENGTH:172
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08386
 SEQUENCE DESCRIPTION:
 GATCCTTGTC AAAATGAGAT TCCAGGTCCT AGAGAGCTGC TTTNAGGGAG TTCCAGGACT 60
 ACTTACTATT GGTGATGCAA TAGGAGAACA GAGACCCGNG GGCTGCTTG GGGGAGGGGG 120
 GAACTCGAGA ATGTATGGAT TTACCTGAAA ACAAATTATT CATTTAATCA AA 172

SEQ ID NO:7150
 SEQUENCE LENGTH:250
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08387

SEQUENCE DESCRIPTION:

5 GATCACCATT CAATTTGAGT TTCCAGGGGG AAGTGCATGT ATAATGAAAT GATAATGGAC 60
 TTCAATGAAG AATGTCATCA ATTATGTACA TATGTATTN CTTTAAATAC AAAGTGTAAT 120
 TTTGTGCCAG TGAATGGAG TCTGAATAGT TATGTGTTT TTTATCCCT GGAAATATTT 180
 ATTAAACTTT ATAGTTTATC CGAGTATGTT GTATGCTTG ACAATAAATG ACTATTTTCT 240
 TCAAAGCAAA 250

SEQ ID NO:7151

10

SEQUENCE LENGTH:282

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08388

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SEQUENCE DESCRIPTION:

20 GATCATCACC ACCACGGTTC CTGAGGACAT CATGTCCAAC CGAGGCAATN ACCTCCTGCT 60
 GAAGCTCCTG CAGNATCGGT ATCCCCGGGT GATGGCAGAG GAGGGTCTTC GAGTGGTGAG 120
 GCAGTGTTG GAGGCCTCCT CACAGCTGGA GGAAGCCTCA NTTATAGCC GATGGGAGGT 180
 GGAAGAGGAC TGGTGTCTGT NTGTCCTCCG CTCCTACCAG GCAGAACACG GGCCCGACTT 240
 CCCCTGGAGC GTGGGGGAGG ACATGAGTGC AGATGGACGG TN 282

SEQ ID NO:7152

25

SEQUENCE LENGTH:391

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08389

SEQUENCE DESCRIPTION:

30 GATCAAAGTA GAAGCNTTTG AATAATGAAT TTAAGGCCAC GCAAAANCTT TCCTCACCTA 60
 TTGACATGAC ATCTAAAAGA CATTTGGAAC TCGAAAAGCA TAGTGCCCA AGTATGGTAC 120
 ATTCTNCTCT NACTCCTTTC TCAGTGCAAG TANCTAACAT TCAAGATTGG NCTCTCAAAT 180
 CGGAGCACTG GCATCAAAAA GANCTGAGTG GCAAAANCTCA GAATAGTTTC AAAACTGGAG 240
 TTGTTGAAAT GAAAGACAGT GGCTACAAAG TTTCTGACCC AGAGAACTTG TATTTGACGC 300
 35 AGGGGNTAGC AAACCTATCT GCAGNGGTTG TCTCACTCAA GAGNCTTNTA GNCACACAAC 360
 CAGTCTCTGT TTCAGACTCT GGGTAANTTN N 391

SEQ ID NO:7153

40

SEQUENCE LENGTH:119

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08390

SEQUENCE DESCRIPTION:

45 GATCTAAAAA TAATTCTTAA TCATGAATTT AATTCACGAC TTCCTTGTTT TCAGTCTTAT 60
 TCAGCTTACT GTAGGAGATT TTAATANACT AAAGCTTTTG TTGATTATG AGTTAGAAA 119

SEQ ID NO:7154

50

SEQUENCE LENGTH:284

SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS08391

SEQUENCE DESCRIPTION:

5 GATCTTGTAC CAAGCCAACG GGTNTCCTGG CTCTCCTGCC CACAGGATGA ACATTTTCGG 60
 CTTCTTAGG AGTTTGGCC TACCGTATTC CAAAGCGTGT GCTGGTTTCT CATATTGTCT 120
 GTAGGCTCAC TCAGCCCGCA GTTTATGTGT GTGCTTTTT CTATGAAAA TGATGTATTT 180
 TGCTACTTCC TGTGTACAAA GTTTATTGT AAATGTTTT TGTGCTTGC ATGAACAGGG 240
 10 GCCACGTTGT TGCAATTGTT TCAGTAGAAC TGGTTTGATT TCTN 284

SEQ ID NO:7155

SEQUENCE LENGTH:289

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08392

SEQUENCE DESCRIPTION:

20 GATCGAGACC ATCCTGGCTA ACACAGTGAA CCCCGTCTCT ACTAAAAATA TAAAAAATTA 60
 GTCGGGCATG GTGGCGGGCG CCTGTAGTCC CAACTACTCG GGAGACTGAG GCAGAAGAAT 120
 GGCCTGAACT CAGGAGGCGG ANTTGCAGTG AGCCAAGGCG ACAGAGCAGG ACTCTGTCTC 180
 AAAAAATAA AAATAGTGCA CTGTCCTTCG AGAAAGTTT CTAACATCTA GTAATTTGTA 240
 ACTTAGAAGT GGAGTTGCCT TGTGGATGTC TTTTTCAT TCTGTAGGN 289

25 SEQ ID NO:7156

SEQUENCE LENGTH:293

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08393

30 SEQUENCE DESCRIPTION:

GATCCAACG CAGAGATGGC AGCTGAGTCA TTGCCAGTNT CCTTCGGGAC ACTGTCCAGC 60
 TGGGAGCTGG AAGCCTGGTA TGAGGACCTG CAAGAGGTCC TGTNTTCAGA TGAATGGG 120
 GGTACCTATG TTTCACCTCC TGAAATNAA GAGGANGNNT CAAAAATCTT CACCACTCTT 180
 35 GACCCTGCTT NTTTGGCTTG GCTGACTGAG GAGGAGCCAG NACCAGCAGA GGTACAAGC 240
 ACCTCCAGA GCCCTCACTC TCCAGATTCC AGTCAGGGCT CCCTGGCTCA TGN 293

SEQ ID NO:7157

SEQUENCE LENGTH:69

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08394

SEQUENCE DESCRIPTION:

45 GATCAGCATG TAAATAATNA AGCATTTTGG TAGGATGTCT ANNGTAATAA ACAATTTTGG 60
 TACCATAAA 69

SEQ ID NO:7158

SEQUENCE LENGTH:312

50 SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS08395

SEQUENCE DESCRIPTION:

5 GATCTTGGCC TCCCAAACCC CTTATTTTAA AAATACTTTG CGCCTTGCTT TGATAATTG 60
 TATTATGTAT CCAAAGTAA ATNATCTGCT TTCTGCATTA GAATGTAAGC CCCCTGAGGG 120
 TTGAGTCAGT CTGTCTTGTT TGCTGTGCCA CGCCTGATGC CCAGCCCAGC AGCATGCTTT 180
 GTACACTGAT ATATTGGGTA AATTTTGTG AATAAATTAA GCTCAACTAT TTGTATTTC 240
 10 ATAGNTTGAG TTGTATTGCT TCCTGTTCTT CAAGCTTAAA TTTGACCTGT CTAATAAAAA 300
 GGAGNAATTA AA 312

SEQ ID NO:7159

SEQUENCE LENGTH:92

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08396

SEQUENCE DESCRIPTION:

20 GATCTTTTGT GGTCTGTAA AAGATACTGA GGAATGTNTT TCAGCCAAGC CAAGAGGATG 60
 GTTCAATAA ACCTAATAAT CTGAAGTCA AA 92

SEQ ID NO:7160

SEQUENCE LENGTH:61

25 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08397

SEQUENCE DESCRIPTION:

30 GATCTTGCAA TATATACTCA GGAAATTACC ATGGGCAGAA TAAATGCTT TAAAAATAA 60
 A 61

SEQ ID NO:7161

SEQUENCE LENGTH:319

35 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08398

SEQUENCE DESCRIPTION:

40 GATCCAAACT GNAGAACTTC TCATTCATCC CCAAGGCCTC CAGGCAGTAT CCAATGGGGA 60
 ATCAGCTCTA AAAGGAACCA GACCAACGTT TTCCAGCCCC TTNATTCTGC TTCCNTCTGT 120
 NTNAGGAAAG GNTAGAAATN TTCAGGGCAT CATCATACAG GCTCCTCATC TACAAAGTTC 180
 CAGTAGCAGT GACGCCTACA CGGAAGANTT GGAAGTCAA ACAGGCTGGG GTCACCTCAN 240
 TNGCATCTGA CGCTGTCAA CCAGAAGTTC GATTTNNTN CTGGGGGTGA AGGTGGAAAC 300
 45 AGACTGTACT TAAAGGNTN 319

SEQ ID NO:7162

SEQUENCE LENGTH:134

50 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

55

EP 0 679 716 A1

CLONE:HUMGS08399

SEQUENCE DESCRIPTION:

5 GATCCANAAA TAAGTTTAAC TTACGGAGGC TTCTAGTTCT AAAGATGCAA NGTAAGATGC 60
CAAAGCAGTG GAAAGATTGA AGTCAAACAG TCGGGCCCAT GTGTGTGTCN TACNTCAACC 120
NTTNGTGTGT TATN 134

SEQ ID NO:7163

SEQUENCE LENGTH:147

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08400

SEQUENCE DESCRIPTION:

15 GATCTATGTG TCTGTNTGTG TGTCATCCC GCCGACCCCC CAGACTAACC TCCAGGCATG 60
GACTGAATCT GGTTCCTCCTC TTGTACACCC CTCAACCCTA TGCAGCCTGG AGTGGGCATC 120
AATAAAATGA ACTGTCGACT GAACAAA 147

SEQ ID NO:7164

20 SEQUENCE LENGTH:320

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08402

SEQUENCE DESCRIPTION:

25 GATCTAGNAA CTAGNAAATA CTATTTGTCC CAGCAATCCC ATTACTGGGT ATATACCCAA 60
AGGAATATAA ACCATTTAAT TATAAGATA CATGCAGATT TTTATTCATT GCAGCACTCT 120
NNACAATAGC AAAGACACAA TAGCAAATGC CCATCAAAGA TAGACTGGAT AAAGAAAATG 180
TGGTACATAT ACACCATGGA ATACTGTGCA GTGCAGCCAT TACAGCTTTT GGTGATACAG 240
30 TGAATCAGAT TTTNNATTAA TTCTTTTAAT TGTTTATTAC TGNNGCTGGA NNAGTAATGG 300
TTTGTATTGG NATCCTTGNN 320

SEQ ID NO:7165

35 SEQUENCE LENGTH:74

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08403

SEQUENCE DESCRIPTION:

40 GATCCATATT TTAAATTGCA ACCCTGTTCC CTATGATACC TATCTGTCTA TGAATGAAAC 60
AAAGGTTTAA CAAA 74

SEQ ID NO:7166

45 SEQUENCE LENGTH:178

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08404

SEQUENCE DESCRIPTION:

50 GATCCAAAGA ACTTCTGTTA TATGGGTTAT AGCTGTTAAT ATTTGTCATA TTCGAAATTG 60

55

EP 0 679 716 A1

AAATAAGATT TTAAGGTATT TTAAATAAC AGTNCAAAAC CATTGTATNT TGACATGAAT 120
AACACTTTTA TGAAATAAC TATATTCCAA AACAATAAAA TTACCAAAAA GTTGCAAA 178

5 SEQ ID NO:7167
SEQUENCE LENGTH:98
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08405
10 SEQUENCE DESCRIPTION:
GATCACAGAT TTCATCAGGA TTGAGCTGCG TGGTAAGCTC GCCACTCTGG ACAAGTCACA 60
GGGTACCCAT TATTAGCAA TAAAAGCTTT AACTCAAA 98

15 SEQ ID NO:7168
SEQUENCE LENGTH:110
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08406
20 SEQUENCE DESCRIPTION:
GATCTCTGTG GCCTAGGTTT TGTACATACA GAAACTGCAT GGTATTTAAA TTATTGTTTG 60
TCTCTGATGA TGTATGCAGT TTCTTTAAAA ACAAAACAAA AAAAAGTAAA 110

25 SEQ ID NO:7169
SEQUENCE LENGTH:298
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08407
30 SEQUENCE DESCRIPTION:
GATCCTGAAG AGCGCTGAGG GACGGGTGCT TCGNCGANCA GCACCAGCGA NACGTCAAAC 60
TCATGAGACA GATGCTAATG GATGCCGNCC TCCCTANTTG TCCACTGCCC CAGCCACATC 120
ATCCCTGTGC GGGTTGCAGA TGTGCTAAA AACACAGAAG TCTGTGGATG AGCTAATGAG 180
CAGCCATAAC ATCTACGTGC AAGCAATCAN TTACCCTACG GTGCCCCGGG NAGAAGAGCT 240
35 CCTTCGGGNT TGCCNCNACC CTNNNGGGG GACCCAGGA TTGNTGGAAC TAANTTCN 298

SEQ ID NO:7170
SEQUENCE LENGTH:329
40 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08408
SEQUENCE DESCRIPTION:
GATCCACCCG CCTCGGCCTC CCAAAGTGCT GGGATTATAG GCGTGACCNC TGCACCCGGC 60
45 CAAAAGTTGA TTTTAAATTA CATAAAATC GTAAAACTT CTAGTAAAAA CTTGATTNNG 120
NGANTACAGT TATATTTTAA AACCTTAAGG TGACAAGCAT TTTCTATGCC TAAATCTTCA 180
TTGGTTTGCC TGGAAAGAGT CTCTGTAAA AGATTTTCCA TATTCAAAGT AAAAGGAAAG 240
ATTTCTTGCT TTCTAATTGT CTTTGGACA CATGCCTATT TTCTTTGAGG TATAANCCTT 300
50 TAGATGTGAA ANATGTAATT TCCATTNN 329

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SEQ ID NO:7171
 SEQUENCE LENGTH:213
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08409
 SEQUENCE DESCRIPTION:
 GATCCAGATT TTTTCTGTTG TGTCTGAAGA GAACCATGAC ATTTAGAAGT ATATTGGTAA 60
 TCTATTAGGT CCATTGGCAA AGTATATTGG TCCATATTCA AACCTATTCC AAAAGGTTAG 120
 10 AAGTGTTTAA GATTCCCTTA TTGTGGTACC ATGTCTGTTT ACCTATGCTT ATGAGCAACA 180
 ATAANTTACT AGTTTACCTT CTGGAATTTT AAA 213

SEQ ID NO:7172
 15 SEQUENCE LENGTH:332
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08410
 SEQUENCE DESCRIPTION:
 20 GATCTGGTGG GCAGTCGAAC CATGGTGAAC TCCACCTCCG TGAATAAAC GGAGATTCAG 60
 CGTGGGCGAC TGAATCCAGC ACGTCTGTGT GAGTANCGGG ACAGTAAACA CTCCACATNN 120
 NNTCAGTTTT TCACTTCTAC CTACATATTT GTATGTTTNN NTGTATAACA GCCTTTTCCT 180
 TCTGGTTCTA ACTGCTGTTA AAATTAATAT ATCATTATCT TTGCTGTTAT TGACAGCGAT 240
 25 ATAATTTTAT TACATATGAT TAGAGGGATG AGACAGACAT TCNNCGGGNT ATTNCTTTTA 300
 ATGGGCACAA AATGGGCCCT TGCCTCTAAN GN 332

SEQ ID NO:7173
 30 SEQUENCE LENGTH:332
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08411
 SEQUENCE DESCRIPTION:
 35 GATCCAGAGG CTCAGTGAAC CCCATCAACT TGGTGGCCCT GGTGTCTCAC ACTTGTATCC 60
 TTCTGCCCTC GAGACCTGGC ACAGCAGTAT CCCTTGAAGA AATCCTGAGG CTTTGTAGAG 120
 TGCTCCTTGA CCATGTTTAA TAATTCTTCC CTCCCCTGCT TGTNTATTTT TTTCTCTTCA 180
 CGGCTCTTCC TATACCTTAG GCCAGTCTCA AGCACTCACT GGAGACCCTT GGGCCTTGGG 240
 CGACCATTGA GTCCTAGTCT CCCTTGTTTG TGCCCTGTGA GGAGGTAGGT CCTTTTNTCC 300
 40 TCCGGCCTAG TAGGGGACCT TGGGTAACAT TN 332

SEQ ID NO:7174
 45 SEQUENCE LENGTH:127
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08412
 SEQUENCE DESCRIPTION:
 50 GATCTGCTTT GCAGAGGAAA TAACTTGTTT TTTCTCATGT TTCATCTTCT TTTTATGTAA 60

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ATATGTAATA CTTTCCTATA TTGCCCTTG AAATNTTGG ATAAAAGATG ATGTTTAAAG 120
TTCCAAA 127

5 SEQ ID NO:7175
SEQUENCE LENGTH:144
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08413
10 SEQUENCE DESCRIPTION:
GATCTGCCCG CCTCAGCTC CCATTGTGA CCATTTTAT ATGTTTATT ACCTATGTGG 60
TTTTCTTTTC TCATAAATG CTTGTTTATA TATTNCCCC ATTTCTACTT ATCTNGTTCT 120
TATTAATNG CCAAAGTGCA TAAA 144

15 SEQ ID NO:7176
SEQUENCE LENGTH:347
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
20 CLONE:HUMGS08414
SEQUENCE DESCRIPTION:
GATCATCTCT AAAATTAATA AAATTAATAT TAAATGTTGG ACCAAGGATT TCAACCATGC 60
AGTATGTTGC TTCTGGGGTC TGCTGTCCTA TTCCAGCCCT AGTGAATAGT TAATGCCTTG 120
25 AACTATTCTC ATCACCACAG GTTGCCAGGC CACCAAATAA CGAAGCCAAT CCAGGATTTG 180
TCCATACAGA AAATGTGTGC GCTCCAGCT AAATCTCAGT TAAGGAGGTT GTTCTATATA 240
CAGCCAAGTA GCAGAAATAT TCATAAATGT CATTTGCAA TATAATAAAC ANTGTAATGA 300
ATGGGTATTT CCTTCTTGT TAATTCTATG TGCTCAATTA AAAGGNN 347

30 SEQ ID NO:7177
SEQUENCE LENGTH:213
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
35 CLONE:HUMGS08415
SEQUENCE DESCRIPTION:
GATCATACTA ATAATTTAAT ATTTGAATTG TATGGAAGTA CAATTCAGTA TCATTTTACA 60
TATGGTATAT TGTGATGCTG TATCATATTT TATGTTACGG TTTATAAGAA AAGCTCCTAG 120
GTATAAATG CTACATAGCA GGAACCTGGT TTTCAATGT TATTATTTCC TACTGTTTTT 180
40 GACGTAACGG CAATAAATTT TGTTTGAACC AAA 213

45 SEQ ID NO:7178
SEQUENCE LENGTH:161
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08416
SEQUENCE DESCRIPTION:
GATCTTAAGA AGAGAATAGA ATCTTTAATT GACCGGGACT ACATGGAAAG AGATAAAGAA 60
50 AATCCAAACC AGTACAACTA TATTGCATAG AATGTTGGCC TTGCAGCATT TGGTGTGATA 120

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EP 0 679 716 A1

TGCAGTAGCC AGTGGATAAA CTAACCTGTT GATTCAATAA A

161

SEQ ID NO:7179

SEQUENCE LENGTH:69

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08417

SEQUENCE DESCRIPTION:

GATCATGTGC AATATTTCTT ACTGTGCCGA GAAGCCACAA TGAGCGAGAT TAAAGCTGTT 60

TAACACAAA

69

SEQ ID NO:7180

SEQUENCE LENGTH:348

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08418

SEQUENCE DESCRIPTION:

GATCTCTGTT CCTTTTGGCT CTAATATGCT ACAACTGTAG GCCAATTATC ACTTTACCAA 60

TAAAGAGTTA GGCCAGATAA GTGAAATTTA ACTTAAGGGC ACACAGCTAA TAAGTAATAG 120

GCCTAAACTG GATTTCCTTA TTCCAAATCC TGTCTTTTCC CCACTATTCC ATTAGACCCC 180

ACAAATGTTA GTTGTGTGTG TGTGTGTGTG TGTTTTTAAT CACTGTAACC GGGTGCATTT 240

TTTTAAGGCA AAATTTCTCC CTTATCTACT GTGATGACTT CAGAAGATAC ANTGGTCCCA 300

GGGGCCAAGT AGAAAGCATT TTAAAGNTT AATCTGAATT AAGCTGAN 348

SEQ ID NO:7181

SEQUENCE LENGTH:132

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08419

SEQUENCE DESCRIPTION:

GATCCACTTC TGTAATTAAG TACNTGGTTG TCTCGTGATG CGTCTACAGT TATTTATAGT 60

TACAACCATT ACCNGNCACT GTAGATACAC ATATAAAAAT AGNTGTTNGG TATCCATAGT 120

TNCGCTTGTT CN

132

SEQ ID NO:7182

SEQUENCE LENGTH:85

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08420

SEQUENCE DESCRIPTION:

GATCTCATCA CTGCACTCTA GCCTAGGTGA CGGAGCGAGA CCCTGCCGCT AAATAAATAA 60

ATAAATAAAT AACATAGTAG TGAATA

85

SEQ ID NO:7183

SEQUENCE LENGTH:224

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08421

SEQUENCE DESCRIPTION:

GATCATGGAC TGGTGCCTTT GCATTCAGAA GGAGAGCTGT CAGCGTANAC CGAATTCAAG 60
ACCAAGGCGT GCTACCTGAG CTGACAGCTT TTTGAAAGCC GAGCTGTTTC TGAACCATGT 120
ACATACATGT TCTGAAACTT TCTCATCATT TTATGAGTAC TGTCATTGA GAGATGACAA 180
TGAAGATTAG ATGAAATTGG AAATAAACCA ACATTGTTTA CAAA 224

SEQ ID NO:7184

SEQUENCE LENGTH:38

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08422

SEQUENCE DESCRIPTION:

GATCTCTGCA GTATTAATAA GATGTACTCC CAATTAAA 38

SEQ ID NO:7185

SEQUENCE LENGTH:67

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08423

SEQUENCE DESCRIPTION:

GATCTTTGGC AGTGCCAGCA GGTAAATTT TGACCTGCTC AATAAAAAAT AAACACTGGC 60
GTTTAAA 67

SEQ ID NO:7186

SEQUENCE LENGTH:117

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08424

SEQUENCE DESCRIPTION:

GATCTTCAAA GAAGAGAATG TAATTTATAC CATGAATATG TTATTTTTTC ATGTTGGAGC 60
ACTGAAATAC TTTCAGTTTG TTTTCTTATT GTTTAAATAA AATTTTAAAT TCTTAAA 117

SEQ ID NO:7187

SEQUENCE LENGTH:231

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08425

SEQUENCE DESCRIPTION:

GATCTTTGTT ATTTTAGGGG GGAAATCACA TGGTATCTAG CATGGGACAA AGCAGCAAAA 60
TAAACCAAAA GGACAATCTC TATTTTATAT AGAAAATAAA CATACTTTNC TGCATCCTGT 120
GTACTTTAAG TATATCTATA TTTNTNANGG GTTCATACTG TGTGANTTT NCCTNATGAN 180
ATAGTCACTT CCCCAGTGTA TTTTAATGCA AATGCATATN CTATAANNTA N 231

EP 0 679 716 A1

5 SEQ ID NO:7188
 SEQUENCE LENGTH:197
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08426
 SEQUENCE DESCRIPTION:
 GATCTCTGTT CTAAGAAAAG GAGTGTGCCT TGCATTTAAA AGGAAATATT GGTTTCTAGG 60
 GAAGGGAGGA GGCTAAATAA TTGATACGGA ATTTTCCTCT TTTGTCTTCT TTTTCTCAC 120
 10 TTAAGAATCC GATACTGGAA GACTGATTGA GAAAAGTTT TAACATGACA TTAAATGTGA 180
 AATTTTAAAA ATTGAAA 197

15 SEQ ID NO:7189
 SEQUENCE LENGTH:115
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08427
 SEQUENCE DESCRIPTION:
 20 GATCAGGAGA CAGGGAGGCC CCGCACATCA CACAGATAAA GTCAGACAAT TGTAATTAAT 60
 ACTTTTGCTG CCTCAAGTTG TTTTAAAT AAAGTACTTT GAAATGCATG AGAAA 115

25 SEQ ID NO:7190
 SEQUENCE LENGTH:329
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08428
 SEQUENCE DESCRIPTION:
 30 GATCCTTGTG GGCATTATTC TAACTGATAC GTAGACACTT ACTTGGAAN NNNNGGACAT 60
 TATATTAAT GAGTGCTATC TGTGAAATTG GTTATATTAG GTGGCTTGAC TAATGTTTT 120
 CCTATAATTG TATATGACT GCATTTTAA AAAAAACCGC ATTTGCCTTT ATGCTAGATT 180
 GTAAAAAATT ATATTAGAAT GCATAAGACA TGTTTNCN TCATATGCTA GACTTTCCT 240
 35 AGCATTTTCGT ATTTCTGTGT TGTCAGTGTG TGATTTTNA ACCGGAATTT GGTTAAAAA 300
 AAATCTGGTG GTAATATATG TGAGAANTN 329

40 SEQ ID NO:7191
 SEQUENCE LENGTH:353
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08429
 SEQUENCE DESCRIPTION:
 45 GATCTTCATT CCAAATTTA CTGCATGGAA ATGACTAATA ATGTGCATAA AAAGTTAACT 60
 ATAAGGATGC TCACTGCAGG GTGGCTTATA TTAACACAAA TTCACAAACA ACTCCACTGT 120
 TCCAGCACAG CAGTGTGTTG TAAATCATGA CATGTTTCTA TGCTCAATTA CTATCCAAT 180
 TTTGAGAATT AGGTGGAGA TATGTAATTN CCAATATGAA AATAAATTC CTATATNCTA 240
 50 TTTAGTAAAA NTAATTCATA AATNTCCATT TTNCAACA CTTTCATGAAT TTNTCCAAA 300

55

ATACANATGT TATCTATCCA TTTNGGAATT AGGANTGGTT NNNATCATTT TGN 353

5 SEQ ID NO:7192
 SEQUENCE LENGTH:352
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08430
 10 SEQUENCE DESCRIPTION:
 GATCTCAGTT TTTCCATCAA AATTATAATA TGCTCATGAA AATAATATTA ATTTGCCTTC 60
 CCTTTGCAAA CACCGGCAGT TGAAAGGAAA AGGACGGGGA ATGTGANNNN AAAGAGACCG 120
 CCTGGAATAA ATGCCCCCT ATGATTCTTT AAGGCAGTGG TTCTCGAGCT TGAATTTTCA 180
 15 TTAGGAAATT CTGTGAGGNG NTTGTAACCA GATTCTCTGG TCTGCCACAT GCACCTATCT 240
 NTNGCTGAAT TGCTTTAATA GAATAATGAG AGCAAGTTTG TCTAACTAAT ACCAACCTGN 300
 CANCTTGAAT ANCAATAAAA TGCAATTGG TACATAAATT ATAATGCTGC NN 352

20 SEQ ID NO:7193
 SEQUENCE LENGTH:348
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08431
 25 SEQUENCE DESCRIPTION:
 GATCCTGGAG AGGAGAAATC CTGCGCTCTG CTCATGCGCA GGTCAACAAC GGGCTGGAAG 60
 TGCGGTTNAG CTCTCTGCCA CTTCTTCAA TGTAAAAAGT AGGCATCAAA GATACGACTT 120
 CTACTATATA CATTTNTNTG TCCTTGAGAA CAGAGAAACG TAACCCTGTA GGGGACTGGT 180
 TACAGAGGAC CTCTGGAGC ATCTGCCCA TTCCTCCAAG TCCAAGCACA GCCATGCTGA 240
 30 GTGGAAGCNA GGCCACCATC TGAGTGGATT GGTCCCAAG NTGTAATGAG GAAAGTGGCC 300
 AGGTGCCGGC ACAGTGCCTC TCCAGAGCC TCAAGAAGGG CCATTNGN 348

35 SEQ ID NO:7194
 SEQUENCE LENGTH:349
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08432
 40 SEQUENCE DESCRIPTION:
 GATCACAAC CTACAGTCTAC CTGGAACACC CCACCCAACA TGCCTGCTGC CTGGGGACAT 60
 GCCAGTTTCA TCAGCTCTCC GCCCTACCTC ACAAGCACCC GAAGCTTGTN TCCAATGTCT 120
 GGACTTTTGT GTTCCATCTG GGCCCCGCAA AGCGATGTGT ATGAAAATTG CTGCCCCATC 180
 AACCACCA CGGAACATTC GACCCACATG GAAAACCAAG CGGGTCGTGT GCAAGGAATA 240
 CTACCCGGGG TTCAACCCGT TTCGCGCTA TATGAACCTG GACATATGGA CTACCACAGC 300
 45 GAATAGGAAT GCAAATTTCC CACTGTCTAG AGACTCGAGT TACTTGTNN 349

50 SEQ ID NO:7195
 SEQUENCE LENGTH:342
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08434

SEQUENCE DESCRIPTION:

5 GATCTGCCTG CCTCGGCCTC CCAAAGTGCT GGGATTACAG GCCTGAGCCA CTGCGCCCAG 60
 ACCAAAAGTT TTTTAAATG GATGTTTAC AGATAGAACT AAATTCTTCA TTTACCTCAC 120
 TTAATTTCTG AAAACACTGT GTCCTCCAAC TGCTCTGTAT TCCCAAAGAT TAAAATTAGC 180
 CTTGGGGAGC AGAGCTGGTT CCTCGTAAAC CCCTGTGAGT GCCCAGACCC TTTGGAGGAG 240
 GTTCTCAGTG GGCTTTTGCC TGCTCCACA GATGTAACCC ACCCTTCCTG CTGCTCCTAA 300
 10 GACCACCATG AGGAAGGCTG GGAATTGGCT CTTGAATTAA NN 342

SEQ ID NO:7196

SEQUENCE LENGTH:364

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08435

SEQUENCE DESCRIPTION:

20 GATCTGCATT TNNAGGTTCC GAGCCTGACC CCTGGAACCTG ACCAGCGCTC GATTGTNAGC 60
 CTTGGCCTGG GGTTTAAACC TTGCCAGTGA AGTTTCGGTT TTAAAGTGAT TAAATGTCAC 120
 TTCTCATCA GTTTCACCTC TGGAGGTTTT CTTATCCTAC TCCCTGGTGC CAGGGACGTA 180
 CCTGGGAGTT TGAATCAGGC CCATTTGAGC GTGGCAGCCG TGTTGGGTGA AGGTCCGGGG 240
 CTCGGTGAGG CACTGGGGGG GTTTTCGGGA GGAAAATGAA AATGCTTCTA GAATGAGTGA 300
 25 ACCACATCAT AGCTCTCACT GTTTTTTCAA TAGCTACTTT TTTTAGCAGA CACAGAGGCC 360
 ACAN 364

SEQ ID NO:7197

SEQUENCE LENGTH:30

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS08436

SEQUENCE DESCRIPTION:

35 GATCTCAAAA TAAATTCAGA AAGAAATAAA 30

SEQ ID NO:7198

SEQUENCE LENGTH:196

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS08437

SEQUENCE DESCRIPTION:

45 GATCTGCCTA GCCAACTTTC TTAACCCAAG CCCAACTGA AAGTCTAAAA GATGTATTTC 60
 TGGGGAATTA CCCACCTCA ACATTCTCAG GTTNCCAGT GTNCTCTGTC TGTNTCAGTA 120
 TTTTAAAAAC ACTAACTCAA TAATNCTCAA AAATCAAAGT CTGCTTTNAT NATCTGGTAA 180
 TAAAATAATA GGCAAA 196

SEQ ID NO:7199

SEQUENCE LENGTH:102

50 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08438

SEQUENCE DESCRIPTION:

5 GATCCGCCCA TCTCCGTCTC CCAAAGTGCT GGGAATNACA GGC GTGACAN CCAAACCCGG 60
CCTGATTAAA GTTAAATAAA TACTAGTTCC CTTCTCGTCA AA 102

SEQ ID NO:7200

SEQUENCE LENGTH:341

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08439

SEQUENCE DESCRIPTION:

15 GATCAGTGCA TTTAGACATC CAGAAGTTAC TTCCACTATT GAGGAAGTAT GTAGACATGA 60
ATTTAAGCTT GTACATGTGC CAATCTTGTT AATGTCACTT GACCACTGAT TCACATAGAC 120
AGACATTAAC GTCTGATGCC TCAGGTTGAT GGC GAACTTN CTTCTAGTG TCTTAACATT 180
TATGANCTTA TGT TTGACTT GAGGGGCATT TCAGCAAAGC TGTGNAGGAA AAGTAGGTTT 240
TGGNCTGATG TCTGCTTTGT NCCATAGACA TTTTGGAGCA TAAGGNAGTA TCTNCTTCTG 300
20 CTATATGTNC AGGGCTTGTC CCCAAATNAA TTCCTGATTT N 341

SEQ ID NO:7201

SEQUENCE LENGTH:110

25 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08440

SEQUENCE DESCRIPTION:

30 GATCTNCTAC AGTTCCTNCA AGCCTAGACA ACCTAAGAAA CTTTACTATC AGCAGCTTAA 60
GATGAAAATC ACAGACTTTA AGAACAGGCG AAGTTTAAAN TGTATATNNN 110

SEQ ID NO:7202

SEQUENCE LENGTH:45

35 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08441

SEQUENCE DESCRIPTION:

40 GATCCAAAGA AATAATCCAA AATAAAAAGT TATATCCAAG ATAAA 45

SEQ ID NO:7203

SEQUENCE LENGTH:202

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS08442

SEQUENCE DESCRIPTION:

GATCCACAGT GGAGTTT TAG TAATTATATT TTNTTGATTT CTTCAATTTT TTTTCTGGTA 60
TAAAAGTAGA GATAATGTGT AGTCACTTCT AATTAGTGA AACCAATTGT AATAATTNTG 120
50 GAAATATTTT GTCTTTAAGT GTAAATATTT AAAAATTTGG CATACCCTAA TGTNAATAAT 180

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AAAAAGAACT ATTTGCAGAA AA

202

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SEQ ID NO:7204

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08443

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SEQUENCE DESCRIPTION:

GATCGGAAAA CTCTGATTGA ACCATTCACT TTATTAGCCA GTAAACTTTC TCAACCCAG 60
 AGCAAATCAA GCTTTCCTTT GCCTGGGAGG AAAANCATAG GGAACCTCTAA TTCTGAGAAC 120
 TAAAAATCAG TAATTTCCAC AATTGTATGT TGAATAGTGA TTGCCTTTAN GNGNCTGTGN 180
 NCATGGAGTA ATATTACTAT TAAAAATAGG CCATTGTAT CTACCTTTGG TCCTTAGGAA 240
 AATTCCTAAG GAAGTCAATT AATGCACTTT NAGATGTAA AAGTATTGG GCTAAGGTTA 300
 TTATTGCCTG ATATGGAAAT AATTATNNTT CTGNTTN 337

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SEQ ID NO:7205

SEQUENCE LENGTH:351

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08444

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SEQUENCE DESCRIPTION:

GATCAGTGAC AATTTACAGG AATGTAGCAG CGATGGAATT CCCTGGAACA GTTTTTTGTT 60
 TTTNTTTTTG TTTTGTGTTT TGTGGGGGGG GGCAACTAAA CAAACACAAA GTATTCTGTG 120
 TCAGGTATTG GGCTGGACAG GGAAGTTGTG TGTGGGGTG GTTTTTTCT CTATTTTTTT 180
 GTTTGTTTCT TGTTTTNAA TAATGTTTAC AATCTGCCTC AATCACTTTG TCTTTTANAA 240
 NGNTTCACCC TCCAGTCCTN TCTCCTCCCC CCTACTCAGG CCCTTGAGGC TATAGGGGGT 300
 TNCTTNGAGG ACCTCANCAA AATCCCAATC CAAGTNAACC TTTTGACAN N 351

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SEQ ID NO:7206

SEQUENCE LENGTH:273

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08445

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SEQUENCE DESCRIPTION:

GATCTATTTT TTTAAAAGAT ATTTGTGCAG AACTGCATG TAACTCTAAG TTTTACTCCT 60
 AACATACATA TGTTGGGGA AGTATTCTAT TCTATACTTG CCAATGTGGA GAACAAAATA 120
 GTTTTTTTAA GAATGAAGAA GTATATATAT CCATTCTGTA TTTTACGTGC AGCAGAATTA 180
 TCTCCGTAG GATTTTTTTG TGTATTCACA AGGTGATATT TGTATTGTAA AACAATAATG 240
 GTGAAGGAAA TAAAAAGGCT TTTAAAAATT AAA 273

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SEQ ID NO:7207

SEQUENCE LENGTH:328

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08446

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SEQUENCE DESCRIPTION:

5 GATCAGATTT TTAATAATG GATTTCAT ATAAATAATA TTATCAAAA AGGATTTCTC 60
 ATATAATAA TATTATCAA AAAGCTGATT TTAAGTTT CTCCCAAAGT CTTATTCTAG 120
 TAATTATAGA GACCTAGGTA ATGAGTGGCA GATATATCTG CCTTCAGAT ATGCCGTAAT 180
 GTGAAAAATA ACACAGTCAT GTGATATTCT TNATTAATA AACTGTGTT GTTTTATTT 240
 TGGAGTAGTT CTCATAATTC ATTGGTAGGG AACTATCCAG TATTTATATT CCTATGTATG 300
 10 TNTATCAGNT TAATTTTGAG GCTTGGAN 328

SEQ ID NO:7208

SEQUENCE LENGTH:145

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08447

SEQUENCE DESCRIPTION:

GATCTGTACA TACAGTGTTA AGAATGTAGA TATTAAAATT GTTATATTTA GCTGTTACAT 60
 20 AATATTAAGA CTCAGAGTTA AGTAATTTCA CTGAAATTGA TTGCTTTTTG TGTCTTGAG 120
 TCAAAATAAA TAACTGAAAT CTAAT 145

SEQ ID NO:7209

SEQUENCE LENGTH:227

25 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08448

SEQUENCE DESCRIPTION:

30 GATCTGTTNA CATAAGCACT AGTCACAGGA CTAAGTGA ATTTATCAA TTTATTCTAT 60
 TGTGTTTATT CATGATATCA CCTTAGTAAT TTTGAAAAA AAAAANGTAC CACACTTGGC 120
 CAAANGCTCA TCAGGCTGAT TGTTANCTG TNGCTGATTG CTTATCTTTA TATTGTATGA 180
 CCCNGGGGNT NGNTTTGAAN GTTATATTAG GTTGAAATG ANTTATN 227

35 SEQ ID NO:7210

SEQUENCE LENGTH:333

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08449

40 SEQUENCE DESCRIPTION:

GATCTTTGAA CTCTGCAATA TGAAGCATAT ATGTGAGTGC TTAGCTTTCA CTCTCTTTCT 60
 GTGTGTGTGT GTGTGTATAC AGATACTAAG ACACAGATAC ATTCACATAT ATACATCTAA 120
 ATATATATAC AGTACAGTCC ATATTCAGAT TTCTCAAGG TTATGGGTCT ATTGTTTAAA 180
 45 TTCGGAATCC AGTCAAGGAC AACACATTAA AATTGATTGT TTTGCCTCTA CAGTGTGTTT 240
 TTTTGGTTTT CATGACATTG ACTTTGATTT GAATAGGTCA GACCAITTTCA TGTACTTTAT 300
 ANAGGCACTT TGTAGATTG TCTGATGATT TGN 333

50 SEQ ID NO:7211

SEQUENCE LENGTH:151

SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear
 CLONE:HUMGS08450
 SEQUENCE DESCRIPTION:
 5 GATCAAGAGA GTTCGATTGG CCAAAGAACA GGAATCCCGG GCAGATTGTA TCAGTGAGTT 60
 TATAGAATGG CAGTATAATG ACAATAACAC TTCTCATTGT TTTAACAAAA TGACCAATCT 120
 GAAATTAGAG GATGCAAGGA GAGAAAAGAA A 151

SEQ ID NO:7212
 10 SEQUENCE LENGTH:275
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08451
 15 SEQUENCE DESCRIPTION:
 GATCTGACAG ACGTCAATAC CTGCAGGCGT CTACAGAGAG GAAATTTGAT AAAAATATTT 60
 TAGTCACTGA CTACACTACC ATTATCTGAA GCATTCAGAT AAGACTTAGC AGAGAATGTC 120
 AGATTTTGTT TTCAATTATA ATTTAGAAA TGCAAAATCT TATTTTCTCT AAGTTGCTTA 180
 TTGTCTATAT AAATGTTTAA TATCATTTAA AAACATGCAG AATAATACAC CTGCAGAATA 240
 20 AAAGAGCTGA AATTAAAATA TCTTTTCAA TNAAG 275

SEQ ID NO:7213
 SEQUENCE LENGTH:103
 25 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08452
 SEQUENCE DESCRIPTION:
 GATCACTTGA GTCCAGGAGT TTGGTGTAC AGTGAGCTAT GATGGCACCA CTGCACTCCA 60
 30 GCCTGGGCCA CAGAGTAAGA ACATGTCTTT AAGAAAAAAA AAN 103

SEQ ID NO:7214
 SEQUENCE LENGTH:144
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08453
 SEQUENCE DESCRIPTION:
 GATCTGTAGA CATGAAGGCA AAGCTCTTGT ATTTTTTTT CATCCAAACA CCTCAATTTA 60
 40 TTTTATAAAT TCGTTCATTT TTCCTGTTAT GTTTATATA ATATATGGAC TAAACAAAAT 120
 AAAATAACAG TGCAAAAGAG GAAA 144

SEQ ID NO:7215
 45 SEQUENCE LENGTH:164
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08454
 SEQUENCE DESCRIPTION:
 50 GATCAGAAGA GGATGACAAT GGCAACAAGT GTTGGAAGT TCCAAGGTGT GTTCAAAGAG 60

55

GCTTGCCTTG AGGGAGGGCT GGAATCTGTC TTCCCTGACT CGGCTCCTCA GGTCTTTAGC 120
CTCCACCTTG TCTAAGCTTT GGTCTATAAA GTGCGCTACA GAAA 164

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SEQ ID NO:7216
SEQUENCE LENGTH:332
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08455

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SEQUENCE DESCRIPTION:

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GATCTGAAGT CCCCCAACC CTCTCCAAGT AAAGTAGTTT ATTAGAAGCT CTTAAGTCTT 60
TAATAGACCT GCATATTTC TTCCCTTATG ATTCCTATA AAATATAGTT TATGGTGTTA 120
TATTTTAAAC TGAATACTG TACATATGTA AATAACTCTT GACAGGAAGA AAATATATTA 180
ATGTTAGTATT TGCCCCCTAT CAGTGAGCTG AACAAATACA TCATTAAAT CTATGCTGCA 240
CTTTGAGTTG CTACAAATAT GGTTCGNTTT GTTTATTTT GAAAAATGTG ATAAAGAAAT 300
CTAAAGANTG NNNNAANTAA NNNNNNNNG NN 332

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SEQ ID NO:7217
SEQUENCE LENGTH:277
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08456

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SEQUENCE DESCRIPTION:

30

GATCTGTACC ATCCTGTTGC TGCTGTATGC GTTCCATTGA TGGGACATCT TCAGGGACTC 60
TTGACAGCCA CCGCTTTCAC ACCCTGGTCT GGAATAAGGA AACATCGGAG GGAGAAGTTG 120
ACTGTCTTGA TAATTAGCCT GACCAGCAGG ATGAATGCAA GACTGACAGT GATGGACTCT 180
GTGACATGGT CAGGTTGAGC TGAAGCCACA GTTCTCTGT GCTGTGTTTT CTAACACATT 240
TTTCTGTTTT TAATTAATAA AANNNNNNNA GGGTAAA 277

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SEQ ID NO:7218
SEQUENCE LENGTH:198
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08457

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SEQUENCE DESCRIPTION:

GATCTTAAAC TCCTGACCTC AAGCAGTCTA CCCCCGTTGG CCTCCCAAAG TGCTGGGATT 60
ACAGGTGTGA GCCACTGCAC CCAGCCAAGT TTATGTATTT TNATTCTTTT TCATGAATGT 120
GATTTGTAA ATNAGTGATT AAATATAATT TTNATATATT TGTATGACTT TNCTGTAAAT 180
AAAATTAAAT ATCAGAAA 198

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SEQ ID NO:7219
SEQUENCE LENGTH:409
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08458

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SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

5 GATCTGCATC TNACGCCAC TGCACACCGT TCCTCTCCAT CTGCCCTTCC CTCCCAGGCC 60
 CCCGGTCGGG GGAGCAGGTG TGGCAAGTGT TAGGAACTCC CTTCTGGTGA CAGCTGGTCT 120
 TCCTAGAGTG TTGCTGGTAA CTATCGCCTC TTGGTCTTCT GACATCATTG CCAGGAACGG 180
 GGCACCTTTT CGTCTGTAG TTTGGCCCTC GGGTTTCCTC ACTAGGTATT GTGTAACCTC 240
 CTCAAAAAA GGTATTGAA ATGCTGAACC TCAGGTTTCA TAGACGTCTT TGTACACTAA 300
 AAATTCTGCA GCAGGAATAT TTTAAACAT TCGCAGTTT TTGTAAGCTA TATTTTTTGG 360
 ATATTAAAT TGCTATNTNA AAANTTTTAA ANTNTTTTT GCTAATNNN 409

10 SEQ ID NO:7220
 SEQUENCE LENGTH:87
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS08459
 SEQUENCE DESCRIPTION:
 GATCTCCTGT TCCGTTAGAA ATACGGTTAT GTTGACATTA TGGTTACTGT GCTTGTGGTT 60
 TTGCAATAAA CATGTATTGA AAGCAAA 87

20 SEQ ID NO:7221
 SEQUENCE LENGTH:130
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08460
 SEQUENCE DESCRIPTION:
 GATCAGATAA AATAACTCCT GACAGAACT GTNNGGAAAC CAGCTGAATG TTTGACCTGA 60
 TGAATGATGT TGTATGGTTT ATGTTAAATG TATATTCTTT TAATCAATGA ATAAAGCATT 120
 30 AAAAATGAAA 130

SEQ ID NO:7222
 SEQUENCE LENGTH:251
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS08461
 SEQUENCE DESCRIPTION:
 GATCAGAAAC ATTTACAATC TATTCTCTTT GAAGCCTGGC ACCTGGAGGC GTCATCTGTA 60
 40 TGAGANNNNN GGACTCCACA ACCTTTTATC ATAACCCAGA CATTCTTTTC TATTGATAAT 120
 AACCAATTGC CAATCAGAAA AATTAAAAAT CTACTTATAA CCCAGAAGCA CTACCCCGCA 180
 ACCCTTGCTT GCTTCAAATT GTTCCAACCT TCTGGACCGA ACCAATGTAT AGCTTAAATG 240
 TATTGATTG N 251

45 SEQ ID NO:7223
 SEQUENCE LENGTH:189
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 50 CLONE:HUMGS08462
 SEQUENCE DESCRIPTION:

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5 GATCGGTGCC ATGGCAAGCA CACCCTCAGT CATCATCCTC ATCAGGACAG NCAGCCNNGA 60
ACTCAAGTCT TACGCTTTGG GAGTTCTTTT TCTCNTCCTT CGTTTGTGG GCTTCATCCC 120
NCCACCCCTC ATCTTCGGGG CTGGCATCGA CTCCACCTGC CTGTTCTGGA GCACGNNNTG 180
TGGGNGAGN 189

10 SEQ ID NO:7224
SEQUENCE LENGTH:97
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08463
SEQUENCE DESCRIPTION:
15 GATCTGATTT GGCCTGGGAC CAGTGTTCAG GTTGGTTTGG TCTTTATTAA AAATCACAAT 60
ATTCCGAAAA CAAAAAACC TAGGAGATAA ATGTAAA 97

20 SEQ ID NO:7225
SEQUENCE LENGTH:255
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08464
SEQUENCE DESCRIPTION:
25 GATCTGAAAT NAGAGGTTTT GGTTCACCT ATCTATGGTA TGTCTTAAAA ATCAACGAAG 60
ATGTCCTTGT CTTTTTNAA TTNCCGAGT GTGTTGCAGT TCCACAGCTC ACTGTTAGGT 120
GGCACATACC CCAAAGTAA AACCTGACCT CGTAGGGCAT GAGTCAAAAG ACAGGTAGGC 180
ACAGGACAGG GCAGTGGTGA CACTACAGCT TTCAGGGTTC CCAGCCTGTC AAGAATGAGC 240
ATGTCTTAGC AGGNN 255

30 SEQ ID NO:7226
SEQUENCE LENGTH:63
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08465
35 SEQUENCE DESCRIPTION:
GATCAAATTT GGAACCTA ACATTGAGAC AATACTATAT TAAAGTATCT CTGGTACCCG 60
AAA 63

40 SEQ ID NO:7227
SEQUENCE LENGTH:55
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08466
45 SEQUENCE DESCRIPTION:
GATCCAGCAA TCTACTGAG TATCTAACCA AAGGAAAATA AATCACTATA CAAA 55

50 SEQ ID NO:7228
SEQUENCE LENGTH:220

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08467

SEQUENCE DESCRIPTION:

GATCCAAGAA CCCTCTCTTG GGGTCTGAAT CATACCCCTT TCGGGTTACA GCCTATCACA 60
GCATTTTAA TCCTATGCAC TTAACACTTC TGTATAGAAA AAGAACATCA CACTAGAATT 120
TTGATGACAA TAGATGACAA TATGATGAAT NTTATTATAT GAAGTGTTAA ACTTATTAGA 180
GCAGATATGG TACTAAAAAT AAAATCCCAC CTNCNCTAAA 220

SEQ ID NO:7229

SEQUENCE LENGTH:114

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08468

SEQUENCE DESCRIPTION:

GATCGCGCCA TTCCACTCCA GCCTACACGA CAGAGTGAGA CTCCATCTCG AAACAAAAA 60
AAGACTTGAT TTTATTTTC ACTATCTTTA GCATAAATTG TTCTAAGAGT GAAA 114

SEQ ID NO:7230

SEQUENCE LENGTH:274

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08469

SEQUENCE DESCRIPTION:

GATCTTATAC AGAGAAGTAT TTTATTAAAA ATTCAAAAGG GAAGACTTTT ATGTGCTCAT 60
TTTGTAATTT TTAATTTTAA ATATCTTTAC ATTGTCTGCC AATTAAAGTG TTTTAACTT 120
GCATTGGAAT GGNCTCTGAA TGTATTTNTN TGGTGTNACG TTATCCGTAG ATTTCTAGCA 180
TGANGTTAGC CTCACGATGC TGTGCAAAGG ATTTTNAAT ATGAGAGTCA CTGANAGAGT 240
TTAANCATCT GTCCATGTTA AATGCTCTAT GGGN 274

SEQ ID NO:7231

SEQUENCE LENGTH:135

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08470

SEQUENCE DESCRIPTION:

GATCCTTTGT TCACGGCTCT CCTTAATGAC TGAGTGAACA GTTCCTATCT GTATATTTGA 60
CTAAACCTTT TCCTAAGCTA TCTCTCATGG TTCCTATGTT TTTTATCAT AATTAAAGC 120
AAAACCATCT GGAAA 135

SEQ ID NO:7232

SEQUENCE LENGTH:238

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08471

SEQUENCE DESCRIPTION:

GATCGAGCGG AAAATTCCAG AACTGTACAA GCCAATATTC AGAGTTGAGA GTCAAAAGAG 60
AACATTGTGC TGTCTGTCAG CATATGTATA TCAGCTACAA AATATATTCA ACTTTGACTT 120
CTTTTGACAA AGGACTTTAG GAAAAAGAGG AACAAAGACA TTATTGAGA ATTAAATTAT 180
ATATTTTAAN TATGACTGTG ACCTTGACTG ATAATAAAGA TGTAATAAGA NTTGCAA 238

SEQ ID NO:7233

SEQUENCE LENGTH:71

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08472

SEQUENCE DESCRIPTION:

GATCAGTTCA GCCAAGCAAC TGACAAATCA AAAACCCACT TGTCAGTTCTN GTAAAATAAT 60
TTGGTCAGAA A 71

SEQ ID NO:7234

SEQUENCE LENGTH:347

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08473

SEQUENCE DESCRIPTION:

GATCTAGAGG GCTTCGAGTG AAGCCAATCA TGTTGAACGA AACTGGGTTT AGTGTAGATG 60
AANTAGAGTG AGGCGTATGC AGTCATAAAA TGAAGGGTGG ACAGGAGATT ACATTGAAG 120
AAAGGATAAG CACTCAGACT TGGTCTCTTT TAAATTAAAA TATTCTTTT CTCATGTGCA 180
AATGTATATC AAAGACTTGA ACAGGTATTC TCTTGAAATT TGAGGATTTT TAGTTTTTAA 240
TGCTGNCTTT GCTTANNATT AANNATGNGT NGCCTATTGG TTCCCAAATT GTNTGGGGNC 300
CCTAANNNGT NCCCTNGAG GGGTNCATTT TTGAAGGNGG GTTANGN 347

SEQ ID NO:7235

SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08474

SEQUENCE DESCRIPTION:

GATCTTGTGT TTGCTATCC TTTTACTGT AAAATGTAAA TATTTTAAGG GATATTTTGA 60
TTCTAAATAT GATAAAATAA TTCTCACCT AAA 93

SEQ ID NO:7236

SEQUENCE LENGTH:417

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08475

SEQUENCE DESCRIPTION:

GATCCCAATT CAACTTATCC AGAATAATTC TCTCCATTCT CAAAATTTC NATGAGGAAC 60
ATNNNNAGAA ACTGTCTCTT GCCAATACCT TTAGTACTGG AATTCTCTCA TCTTCCTCA 120

EP 0 679 716 A1

5 CAGTCTTGGC AGCTAGGAAA AAACAGCTGA TTGCAATACA ACTCAAGTAT TTTGGATGAG 180
 CCTTTACGGT AGCTAAAAAC CCATCCAAAA GACTGCTAGC CAGAGCAAAT GTTCTGGGT 240
 AAAGGTTGAA TTGGTACTTG AGTTTGGCCA GCCATTGANT TACTTCATCT CTCTGGGATG 300
 GNGAACATTC TGATTGAAG GCATTTCCG CACATTCAC TCCACATCT GTGNCCTCCCT 360
 AGTGATTGCC TTTTCCACCA GGAANGCCAA TCTCTGGTTT TCCAAAGGCC CTNGGAN 417

10 SEQ ID NO:7237
 SEQUENCE LENGTH:80
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08476
 SEQUENCE DESCRIPTION:
 15 GATCAGACCC TCCTGTGGGC AGGGTTCTTA GTGGATGAGT TACTGGGAAG AATCAGAGAT 60
 AAAAACCAAC CCAAATCAAA 80

20 SEQ ID NO:7238
 SEQUENCE LENGTH:409
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08477
 SEQUENCE DESCRIPTION:
 25 GATCTGCCTG CCTCGGCCTC CCAAAGTGCT GGGATTACAG GCGTGAGCAT TTGCGCCTGG 60
 CCAGAAAAAT ACCTTTAAAA AGCTGGGCCT GTTGGTGTAC CTAAAGGATA ATACAGATGT 120
 TCCTCTCAAC TTAGCATGGG GGTGTGTCT CAATAAACCC ATTTTAAAGT TGAAANCGTT 180
 ATGTGGAAAA TGCATTTTAT TATACCTAAC CTACTAGACC TCGTAGTTTA GCCTCCCTTA 240
 CCCTACAGGT GCTCAGAACA CTTTNGNAGG CCCTACAGTT NGGGCAGTCA TCTAACACAA 300
 30 AACCTATTTT ATAGGTAAAA GTNGTTTGAA TATCTCAAAG NTAATTTATT TGAATGGITN 360
 CCACTTGAAT NGCGGANTCN ANGTGTTTNN NAACNGTNT GGTNNANN 409

35 SEQ ID NO:7239
 SEQUENCE LENGTH:123
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08478
 SEQUENCE DESCRIPTION:
 40 GATCTCATAG ACTTAGTACA AAAAAAGGTG AAATGTCTTG ATATTTTGT ATATTGATTA 60
 TTGAAATAAC CTTTATGTT GGGCTAAATA CAAATATGTT ATTAATAAATNA ATTTAACAAAC 120
 AAA 123

45 SEQ ID NO:7240
 SEQUENCE LENGTH:265
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08479
 50 SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

5 GATCTGAGGG GGAGAGGGAC GGGTCGGGCC AGCTGGTCAG GAAGGGCCTT GAGGGCCTGC 60
AGCCAAGAGT AGCAGITTTT TCCCCGTCT TGTGTTCTT GGTCCCTGGA CCACAGGGCA 120
CACAGGCTG GACACCATAA GGCTGGTGGG CTTTCAGAAT TGTGTTAGGG NNNNNNGAGT 180
GGCAGGTTCC TGAATCTCGG TCAATATAGT AACCAGCAGG ACAAGAGGTT CAGGAGGAGC 240
CCACATCAGA GGCTTCTAGG GCAAN 265

10 SEQ ID NO:7241
SEQUENCE LENGTH:265
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08480
SEQUENCE DESCRIPTION:
15 GATCTCTTGA ATTGCCATCT GTCATCTAAG TCTAAGGCTT TATGCTTAAT GANTTGTGGT 60
GTACTAGCTG GACTAGCTTC AGGAATGTCA GTGTGTTCTA CCATTAAGGT CAGATTGTCT 120
ACTACATCGA GATGGTGGTT GTAGTTACAG CTACTTTTAG AAACATGTCT ATTTTTTAAA 180
GACATAACAC ATGAATGAAG AAATTCAGAA TTTGGAAAAA AGGTCCGTAA TGCCTGACAA 240
20 AGAAAAATGT TCTCCTGAGG GTCTN 265

25 SEQ ID NO:7242
SEQUENCE LENGTH:138
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08481
SEQUENCE DESCRIPTION:
GATCCCAGCA CCTTCACATC TTCAGCAACC CAGNNGTCA AATTTGACCT GGTGTTATTT 60
30 TAGCCCCAAA TTTATGACAT TACACAATAT TAAATGTAA ATGTTTCTTC ACCCAAATA 120
CTTCTAGATA TNCTNGNN 138

35 SEQ ID NO:7243
SEQUENCE LENGTH:273
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08482
SEQUENCE DESCRIPTION:
40 GATCCTTTGA GAGATGGTTT GACTTTCCCC GAGCAAAGAG CCTGCGTTGA AAAGCGGGGG 60
TGGAATTCAG TCCTCACAGA TAATGAGGGG ACAANACCTA ATTGANCCGN GTATTGCCGG 120
GAAGGAAAAG GCAACGGGCC AAGCCTTTGA CAGGGTGCGA AACTGACTTT NATCATCGTT 180
ATAGTCTTTA AATCCTGGGA AACGAGTTGG CAACCCCAA ATAAGAAGT GTAATGACGT 240
45 CTGATGACTT CACCCAAATA CAGACCATTC CAN 273

50 SEQ ID NO:7244
SEQUENCE LENGTH:132
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08483

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SEQUENCE DESCRIPTION:

GATCTATAAA ACAGACAGAG ATAAGTACAA CAGAATATCT CGGGAATGGA CTCAGAAGTA 60
 TGCCATGTGA TGCTACCTTA AAGTCAGAAT AACCTGCATT ATAGCTGGAA TAAACTTTAA 120
 ATTACTGTTA AA 132

SEQ ID NO:7245

SEQUENCE LENGTH:321

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08484

SEQUENCE DESCRIPTION:

GATCCGCCTG CTTACGCCTC CCAAAGTGCT GGGATTAGAG GCGTGAGCAT TCATGTCCAG 60
 CCAGCACTGT GCTGTTTTTG ACATGCATGT TCTCATTCAN TGCCATGCGG CAGATTCTAG 120
 GATTATNATT ATNCTCACTT TACAGATGAG GGAGTGAGAC TTAAGGGGT TAGGTAATTT 180
 GCCCACATCC CCCAACTAGC AAATGGTAGA GCCAGGATTG GAATTTGTTT AGGCTCTTAA 240
 CCATTGCACT CTTTCAAAAT GTTAAAAATA ATGTTGTGTC TCATTTTAAA AAATTAAATA 300
 AAANCTCTTT GGACCAAGAA A 321

SEQ ID NO:7246

SEQUENCE LENGTH:108

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08485

SEQUENCE DESCRIPTION:

GATCTCCCCA TTTGGAGACA GAGCCATTG GATATTTNCC CCTTNAACTT CTCCANNNCC 60
 TGAAGCGTTC CTTCCTGGA GGAAGTCTTT GGTTCAGGG CTAAACTN 108

SEQ ID NO:7247

SEQUENCE LENGTH:139

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08486

SEQUENCE DESCRIPTION:

GATCTCAGCT CAAATATTCC TTATGCAGAG ATGCCTTCCC CTGATTCCCT AAAGTAGACT 60
 CGGACCCCTC TGTCTGCAT CTCGTGGCA TATCATCTGC CCCTTTTGTA AAAGTTATAA 120
 ACTTGTGGTT ACTTGTA 139

SEQ ID NO:7248

SEQUENCE LENGTH:244

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08487

SEQUENCE DESCRIPTION:

GATCATGTGA GCTCAGGAGT TCGAGACCAG CCTGGGCAAC ACAGCAAGAC CCTGTCTCTA 60
 TTTTATTCTA TGTTTTAAAA TTTAAAAATT AAAAAGAAAA AAGAATAACA TTTTATAAGA 120

EP 0 679 716 A1

AAATATTTTA TGAGAGACTT ATAAAAGAGA CATTAAGAAT ATGAAGTAAT ATATTNCGTA 180
TTCCTTATCT CAATAATGCT GTAGGGAAAA ANCAATTGAA ATAAAAGCAG TAAATGAGTG 240
GAAA 244

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SEQ ID NO:7249
SEQUENCE LENGTH:155
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08488
SEQUENCE DESCRIPTION:
GATCTAAGT GCAGTGTATT TTCTAGGCTG GAAAGTGGAA AATNAAATAT ATTATAATCT 60
TAGGTTACAT AAAGTTTCTA AAGTTTCAAA GAGTCTTGAT ACAAATCAG TTTATATTCT 120
GAAAAATTTT ATAATAAAGT ATTCTAATTT CTAAT 155

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SEQ ID NO:7250
SEQUENCE LENGTH:96
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08489
SEQUENCE DESCRIPTION:
GATCGCACTG GAAGATTTTA TAAAAGAATT TTTGTGGGTC GGGGGGACAG TAACTTCCT 60
GGGCCACGTG GGTCCCTCAG GAGTTTTTCA GGCAAA 96

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SEQ ID NO:7251
SEQUENCE LENGTH:118
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08490
SEQUENCE DESCRIPTION:
GATCTNATAT TGAACCTAT GCCAAATAGC GTGGCATGAG GCCAGAGAAC AGTGCTTNCC 60
TTCCCCGCT CCACCCAAAT TGTCCCTTGC TCTNGATATT ATGCCAGTNC CATATTTN 118

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SEQ ID NO:7252
SEQUENCE LENGTH:391
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08491
SEQUENCE DESCRIPTION:
GATCTGGAAA CTATTGGNT TTTNTTTTCA ACTTTTCATT TGGATGTTTG GCGTTGCACA 60
CACACATCCA CCGGTGGAAG AGACGCCCGG TGAAAACACC TGTTTGCTTT CTAAGCCAGT 120
GAGGTTGAGG TGAGAGGTTT GCCAGAGTTT GTNTACCTCT GGGTATCCCT TTGTNTGGGA 180
TAAAAAAAT CAAACCAGAA GGCGGGATGG AATGGATGCA CCGCAAATAA TGCATTTCCT 240
GAGTTTNCCT GTTAAAAAAA ANTTTTTTTA AGTAAGAAAA AAAAAAGTAA TANCATGGCC 300
AATTTGTTAC ATAAATGNC TTTCTGTGTA TANNTTATTC CTAANATC CTGTTTATAT 360
AANNNTTCAG TAGNTGGAGA AACTTTCCAT N 391

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EP 0 679 716 A1

SEQ ID NO:7253

SEQUENCE LENGTH:385

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08492

SEQUENCE DESCRIPTION:

GATCATGAAT CTCTAATAAA TTAAATCTC TTAAACCAGT AGGTGCTTAA TATTTTTTAA 60
TTTGATTAAT GCCCATTTAA ATCTCATGGG TTCTATTAAA AATATATATA TATAGGGCCC 120
CAATCCATTG CCATCAAATT GCCCTGGAC TTTTCCAAGG TATATTATGG GGTTTTATGC 180
AAAATTCCAA GCTACCATGT AACTTTTTTN AACCATTTAA CAAGGAGGGG GAACTGTTTC 240
CTACCTTCTT TACATGTTGT GCATTGTTGT GGTCCAGAAA TGCCAAACCT TTTTAAAGAT 300
GGTGCAACTT TGAGTCCTTG GCTTGACTAT ACAGGNCCTG AACTTCATGG CATATCAACT 360
TTGNCATATC TGCAGGTGNG CTGTN 385

SEQ ID NO:7254

SEQUENCE LENGTH:414

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08493

SEQUENCE DESCRIPTION:

GATCACCATT TTTTCCAGG CTTTCTGCCT CCNAGATGTG GCACCATAGT GCGGTGCCCT 60
GTGGCTTCAC CGCCCTACTT CCACCTCCGC CCAGCCTGTA ATGTTTATAT AAGCAGCCTC 120
AAGGACCAAG AACCATCTGC GAAAGGACAC ACACAGGAAA TTCATAAAAG AAATCTGANT 180
GGNTAAAACC ATGAAAAAAA GTATGCTTCA TTAGTAATTA AAGAAAAGGCA AATAGAGCTG 240
GAAGCNTTTT NCCNTTNGCA AACCATTAACA GAAAANNNTA NGACCCAATA TTGGCAAAGA 300
GACTACTGAA AANCCATTCC CATACATTGC GTGTGGGNGT ATACATCGGT GCAGGCTTCC 360
TGGATGACAG TTGGGTGATA TGTGTCATGT GGCCTAANAG CCTCCATGTC ANTN 414

SEQ ID NO:7255

SEQUENCE LENGTH:207

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08494

SEQUENCE DESCRIPTION:

GATCCTGGTT TTGTGATTGA CAATATTTCA TGCTCTGTAG TGAGAGGAGA TTTCCGAAAC 60
TCTGTTGCTA GTTCATTCTG CAGCAAATAA TTATNATGTC TGATGTTGAC TCATTGCAGT 120
TTAAACATTT CTNCTTGTTT GCATCTTAGT AGAAATGGAA AATAACCACT CCTGGTCGTC 180
TTTTCATAAA TTNCAATATT TTTGAAA 207

SEQ ID NO:7256

SEQUENCE LENGTH:412

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08495

SEQUENCE DESCRIPTION:

5 GATCAAGAGT ACACTTCCTG GCAGTGATTA AGGAGTGTGT ATCTAACAGA AAAAATATAT 60
 ATACCCTGTG AACCCGAATA TGGAAATTCAG ATTGTTTCTG CCCTCAGTAT CATACTTAAA 120
 AAACAAGCAT ACAAACAAAC ATAAGGGAAC AAACAGCAAC CATAACAAAA ACAAACTTT 180
 AAAGGTGGGT TTTTGCTGTG ATAAATGAAT ACGGTACTCT GAAGGAGAAA AAAGTTTCTC 240
 AAATGAGCTT AAACGCAAG TGATTAAAA NNTAGAGAAT ATAATCTTA ANGCTATTGA 300
 AAGTTTCAAC CAGAAANCCT CANGTGAATT TTGTATGTAA ATGGCATCTT GNNTGTAAGT 360
 10 TCTGTGNTTC TTTAGGCAAC CAATTAGCTG GAACCTTGGT TTTGTGTGGT TN 412

SEQ ID NO:7257

SEQUENCE LENGTH:406

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08496

SEQUENCE DESCRIPTION:

20 GATCTGCAAA TCAAGCTACA TGTATTTGTG TATAAGACCC TGTGTTGAGG ACTGGGTGAC 60
 TTTTCTAAGA AATATGGGGT TTAGAATGGG GTTGACTGTA TTTNTAAACC TAATTCTGGA 120
 GAGAAGATTG TATTNTAAC AGTTTTTGGG GTTTGGCTTC CTTCTCACAT TTCTTTAGCT 180
 TTGAATTTT ACTAAATAAA TTTCTCCTG ATTAATTTT TTTTCCCCA TCTGGGAATT 240
 TGAATCTCG GTGCTTACTG TNACACCAAT TTGTCCAANG GGTGAAATC ACTTAAATGC 300
 CAGACCATGG TAATTTGAG CCATTTGAGG CAGGTGGTGG NCCTTTTNA TACCATCGTT 360
 25 ACGGGGTCCC NTTAAATATT CTGGGGGGTG ATGTAAATNT AACGGN 406

SEQ ID NO:7258

SEQUENCE LENGTH:116

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS08497

SEQUENCE DESCRIPTION:

35 GATCTCCATT TCCGCCAGCT GCCTGTAGCC ACGTGGCATC CTGCCTGTGG TCTGGGTGAG 60
 ATTTACTGTG ACCAGATGTA GAATAAATGT GTCTCATCCT GCATTTTTTT TCTAAA 116

SEQ ID NO:7259

SEQUENCE LENGTH:435

SEQUENCE TYPE:nucleic acid

40 TOPOLOGY:linear

CLONE:HUMGS08498

SEQUENCE DESCRIPTION:

45 GATCACTTTT TAGTAGTTT GAAACCTGCT TTTAGTTTAT ATGTAGTATT TCTACATATT 60
 GTTACTTAAA TTGAACAAAC AATTACTTT TNAATTTTNC TTCTTCCTTT TAAAAAAGA 120
 GTATATAATC TGACAATGTT AGTACATCTT TTAGAAGGTC ACTATAAAGT TGCTTCCCT 180
 ATTTATTTGG AAAGGATTAT TTCCTTATNA TTTTAAAAA TCTTTTAATT NNTGTCTTNT 240
 CAAGTAATTT TATCATTGTA CCAAGCCTAA GGATGAGTGG CAATTTAAAA GNCACAAGGT 300
 50 GTGCATCTTC TATCTGCAAA TACTCCAAAC AGAANTTATT CCAGTTTGTG GATACTTTGA 360
 GTGGCCAGG GAAAAATGTG ATGGTTTTAG TGTNCTGAAT TGNCATTCCA AAGGGTGC GG 420

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AAGGTTNCTG TTAAA

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SEQ ID NO:7260

SEQUENCE LENGTH:62

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08499

SEQUENCE DESCRIPTION:

GATCAAAAAA GGAATGGATA TACAAAGTGT TTTGTGAAAT AAAAGCTCCC TAAAATGGTA 60
AA 62

SEQ ID NO:7261

SEQUENCE LENGTH:389

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08500

SEQUENCE DESCRIPTION:

GATCGCACCA CTGCACTCCA GCCTGGGCGA CAGAACGAGA CCCTGTCTCC AAAGGAAAAA 60
CAAAAAAGAA GAATAAAATA ATTTGGATGA AAATCATGTT TATTAAATA GTAATGTCAT 120
GAGACTATTA AAGATGTGCC AGAGTTTCAA TGAAAATCAT TAAAGTAGGA CAGCTAAGAA 180
ATTAATATTA ATATCAAANT TATTGATANT CTAAATTAT TGATTATTCC TTAACGCACT 240
CCATTCTCCT TTTACATTTT ATCATGTTTC TTTTGAATAT ATGANTTGGC AAAGGCCTTG 300
ATGAAACTGA GTACTAAGAT TTGGTCCAGN GTATGTCAGG ANGNCACCTC AGATTGCCAT 360
TTTAAATAAN GTTGTCATG ACCAATAAA 389

SEQ ID NO:7262

SEQUENCE LENGTH:302

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08501

SEQUENCE DESCRIPTION:

GATCCCGTGC TACATCTCCT CCCCGGACGA GTGCCTCTGG ATGGACTGGG TCACAGAGAA 60
GAACATCAAC GGGCACCAGG CCAAGTTCTT CGCCTGNATC AAGAGAAGTG ACGGCTCCTG 120
TGCGTGGTAC CGCGGCGCGN CGCCNCCCAA GCAGGAGTTT CTCGACATCG AGGACCCATA 180
AGCAGGCCTC CAACGCCCTT GTGGCCAACT GCAAAAAAG CCTCCAAGGG TTTGACTGG 240
TCCAGCTCTG ACATCCCTTC CTGGAACAG CATGAATAAA AACTCATCC CATGGGTCCA 300
AA 302

SEQ ID NO:7263

SEQUENCE LENGTH:297

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08502

SEQUENCE DESCRIPTION:

GATCGGAAGG TGGTGAATCT TTCCAGAGCA GCTGAAAATN TCAGCATGGA GACAGTAAGA 60

EP 0 679 716 A1

AAAGAGACGG GGTGTGGATA AGACTCTGCC ACCGTGTCAC ACTAGCATAG GAGGCTGCAC 120
 GTTCATTGT TGTGTTTTT TTTTCCTTG CCAACCTCCG TTCTATTTAT GTGCAAGCAG 180
 TTTGGATTCA AGTTCTTGTA TCTGTCTGTT CTGGGACCTG GGGATTGTNA GGGTTCCCTC 240
 ACAGCCAGCA CGACCCCGAG AAAGAGGCGT CCCACAATAA ACACGTCACC TGCTAAA 297

SEQ ID NO:7264

SEQUENCE LENGTH:116

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08503

SEQUENCE DESCRIPTION:

GATCAACAGT TCTATTACAG ATTAGAATGT TTCACTTCAC AGTTCTTTGT AAATCATGTC 60
 AATTTTATN ATAGNTGTCA ATTTGGGAAA ATCTGTATTA AAATTTTAA ATGAAA 116

SEQ ID NO:7265

SEQUENCE LENGTH:442

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08504

SEQUENCE DESCRIPTION:

GATCCCTAAT TGGCCAGGCC TGACCCTCTT GGACCTTTCT TCTTTGCCGA CAACCACTGC 60
 CCAGCAGCCT CTGGGACCTC GGGGTCCCAG GGAACCCAGT CCAGCCTCCT GGCTGTTGAC 120
 TTCCCATTCG TCTTGGAGCC ACCAATCAAA GAGATTCAAA GAGATTCTCG CAGGCCAGAG 180
 GCGGAACACA CCTTTATGGC TGGGGCTCTC CGTGGTGTTC TGGACCCAGC CCCTGGAGAC 240
 ANCATTCACT TTTACTGCTT TGTAGTGACT CGTGTCTCC AACCTGTCTT CCTGAAAAAC 300
 CAAGGCCCN C TTCCCCACC TCTTCCATGG GGTGAGACTT GAGCAGAACA GGGGCTTCNC 360
 AAGTTGCCAG AAAGACTGTC TGGGTGAGAA GCATGGCAGA GTTTTCCCA GGACAGGTGT 420
 TGACCAGGGC TTNTTGTITA TN 442

SEQ ID NO:7266

SEQUENCE LENGTH:148

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08505

SEQUENCE DESCRIPTION:

GATCTTCTGA CATTACTGCT GTCTGAGATT TGTATATGTG TAAATACAAG TTCCTTGATA 60
 CCCTAAAACC TTGGATTAAA CAGAATGTGC ATTGTACATC TTAAACAAA ATGTATATTA 120
 ATTTATATA TCTAGTTGTC ACTTTAAA 148

SEQ ID NO:7267

SEQUENCE LENGTH:169

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08506

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

GATCTTAATT CTGTGTCTTA ACAGCAGGAG ATGAAATTCC CACATGGGAG ATGGCCTCCC 60
TGTCACAGGA AGGAATANCA TTCTCCCTAT CAATNNNGAG AAATTGGAAC TGNGAGAATT 120
CTGTATAGAC CCTTGTTCAG GTAATTNTTC TAGTTTAAAN CNTCCCAAN 169

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SEQ ID NO:7268
SEQUENCE LENGTH:89
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08507
SEQUENCE DESCRIPTION:
GATCAAAGAG TCTCATCAA GCAAGTTCTT CCCTCCCTGG TCCCCTTTGG CTTTCTGAAA 60
TNATTAATTA AAAGAAAGGC CCTAGGAAA 89

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SEQ ID NO:7269
SEQUENCE LENGTH:225
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08508
SEQUENCE DESCRIPTION:
GATCATTGNT TTATTTTGTG GTCCACCTTT CAATCACTTC ATGTATATAA AAAGGGACTT 60
TGAATGTAAT AGTCTTCATC TTATTTATTG AAAAGATGTA CTAAAGGAAT GACTATTTTT 120
AAAAAANCT CTAGTATTTA TCAATATGAC TCTAAGGGAA AAATCAAGAA ANTTAGAGGA 180
AGGTTC AANT TAGTAGTAAG GAATAGGANC NNNCATATNA NCTNN 225

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SEQ ID NO:7270
SEQUENCE LENGTH:259
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08509

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SEQUENCE DESCRIPTION:
GATCTGTAGT GATGGAGTTC CTTCTGGTGT CAGCCCCACA GGAGGCTCCC AGGCCTCCCT 60
CACTTCCCAT ACCCAGTCTA GGAGCTCCTT CTGGCTCCCA AGCACCAGAA GCTTTCCTCC 120
GCCTTTNAGT TTTGGTTCCT CCACTGGAAT GTAGGCTCCT CACGGGCGAT GGCTGTCTTT 180
TCTTGACTTT GTATCTTCAC TGCCAAGCAA AAAGTCTGCC AAGTGGGAAT NTTAATAAAA 240
TATTCATTGA ATAATGAAA 259

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SEQ ID NO:7271
SEQUENCE LENGTH:357
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08510
SEQUENCE DESCRIPTION:
GATCCCAGAA GGAATGCTGA CCCCTCGTCG TATGACCTGT GCATAGTCTC CAGAGCTTCA 60
AAGGCAACAC AAGCTCGCAA CTCTAAGATT TTNTNAAACC AAAAAACCC TGGTTAGCCA 120
TCTCATGCTC AGCCTTATCA CTTCCCTCCC TTTAGAAACT CTCTCCCTGC TGTATATNAA 180

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EP 0 679 716 A1

AGGGAGCAGG TGGAGAGTCA TTTNCCTTCG TCCTGCATGT CTCTAACATT AATAGAAGGC 240
ATGGCTCCTG CTGCAACCGC TGTGAATGCT GCTGAGAACC TCCCTCTATG GGGATGGCTA 300
TTTNATTTTT GAGANGGAAA AAAAAAGTCA TGTATATATN CACATAANGG CATANAN 357

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SEQ ID NO:7272

SEQUENCE LENGTH:265

SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS08511

SEQUENCE DESCRIPTION:

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GATCTAAATN AACAGCATTT TTTTCCTTAG CCTCTGTTTG CCACTCTGGG TATCTCTCCT 60
ATGGGCAAAG CCATTAGAAA TGCATAAAAC CTCGAGACAT GGTTTTGGC AAAAAGTCCA 120
TGACTTTAAA CTAGCTCTTT NACTACTGAC CTTTCACAGA GAAAAAATAT TTCCTTGAA 180
AAAAAGTGGG CTTGTNATTT TTNCCCTTGT AGCTTTAAGC AGAGACATAA GTGCCTTGCA 240
TTACACATAG TAAACTTTCT TTAAA 265

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SEQ ID NO:7273

SEQUENCE LENGTH:164

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08512

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SEQUENCE DESCRIPTION:

GATCTTCTCT TTTTTGGGT TCATTTTGTG CTGGGTTTG GTTTCTTCA CAATCTTGAA 60
CATTTTACAG TAGAACTCAT CTAAAAATGG ATTGGGGAT GGGGAAACAT GCACAAAATC 120
TTTTCATAAT TAAAAAGAGC CTTACTTTCT TTACATACCA CAAA 164

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SEQ ID NO:7274

SEQUENCE LENGTH:243

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS08513

SEQUENCE DESCRIPTION:

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GATCCGTTCC GTGGTCGCCC TGCACAACCT CATCAACAAC AAGATTGCCA ACCGGGATGC 60
AGAGANGAAA GAAGGGCAGG AGAAAGAAGA GAGCAAAAAG GATAGGAAAG AGGACAAGGA 120
GAAAGATAAA GATAAGGAAA AGAGTGATGT AAAGAAAGAG GAGAAAAAGG AGAAANAGTA 180
AAACATGTAT TAANTAGCTT TTTTAATTG TAANTTAANA TCTTACANAC TNGAANCNAN 240
NNN 243

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SEQ ID NO:7275

SEQUENCE LENGTH:265

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08514

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SEQUENCE DESCRIPTION:

GATCACAGGC CTTGAGAGAG NAGAATAAGA CACTCTTGGG GACAGAAAGT TAAAACACCG 60

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EP 0 679 716 A1

5 TGTGTCCTCA CTTATATGTG GCGCTAAAA AAGTTGATGT CATAGAAGTA TAAAGTAGAA 120
CAGAGGATAT GGAAGGTGG GAAGGGTAGG GGAAGGGAA GGATAGGGAG AGGTTTGTTA 180
AAGGATACAA AATTACCACT CGATAGGAGG AGTAAGTNCT AAGTNCTAGT ATTCTATACC 240
ACTGTAGGAT GACTATAGTT AACAN 265

10 SEQ ID NO:7276
SEQUENCE LENGTH:261
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08515
SEQUENCE DESCRIPTION:
15 GATCGCCCGC ATTCCGGAGC CTTCAAGTAA ACGGCGCGCC CCACGAGACC CCGGCTTCCT 60
TTCCCAAGCC TTCGGGCGTC TGTGTGCGCT CTGTGGATGC CAGGGCCGAC CAGAGGAGCC 120
TTTTTAAAGN ACATGTTTTT ATACAAAATA AGAACAAGGA TTTTAATTTT NTTAAGTATT 180
TATTTATGTA CTTTNATTTT ACACAGAAAC ACTGCCTTTT NATTTATATG TACTGTTTTA 240
TCTGGCCCCA GGTAGAAACT N 261

20 SEQ ID NO:7277
SEQUENCE LENGTH:260
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS08516
SEQUENCE DESCRIPTION:
GATCCACATC TAGTGAAATG TCAGTGTCAG AATATTATAG ATTATAGCTA AAATCCAGAT 60
TAATACTCAT TTGGGGTTTT TAATAGTGGA ACTTCATAGT AATACAAAAA GCAGATTGTC 120
TTCCTGTCTC CGCTGCTCCC ACAGTAGGTA TTGAAACTGG TAAAANCAGT TTTTGTATAG 180
30 TGTGTGTATA TAAGANANNN TAGATACACA CATTCTTTTT CCTCAGTCAN CACATTGATT 240
GACCACTCTG GCAAAGATGN 260

35 SEQ ID NO:7278
SEQUENCE LENGTH:60
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08517
40 SEQUENCE DESCRIPTION:
GATCCCACTC CACCCACCT CAACTATTTT AACTTCCTAA TTAAATCAGA CTGTTTGAAA 60

45 SEQ ID NO:7279
SEQUENCE LENGTH:156
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08518
SEQUENCE DESCRIPTION:
50 GATCTGAGGA CTAAAAAGA AAACAAAAC AAATGGCAAG TTTCACCTAA GGGTGGTTTG 60
CCCTTAAGAA GAAAGCTGTT GGGACAAAGA CACCGAGCCA TTATACCCAG AATAAAATAA 120

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TACATTTATG CTGGATTTTA TTCAGACCAA ACTAAA

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SEQ ID NO:7280

SEQUENCE LENGTH:329

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08519

SEQUENCE DESCRIPTION:

GATCATTGTT TAGAATATTT CCGCCACCTT AGAAAGTTTT GTCATGCCTT TTTTTCAGT 60
 CAGTCCCTCC CCTTCACTCA CTGCAAGACC ACCACTGTTG TGACTTTTAG CATCTAAATG 120
 TAGCTTTGGA ATCATGTAGT ATGTACTCTC ATGTCTCTGA AATTCATCCA TGTGTTGCA 180
 TAATTCAGTA GTTCACTACT TTTTAATTTT TGATTAGTAT TTCATTGTAT GGGTGTACTG 240
 TAATTTGTTT AACCATTAC CTGCTGTTTA ATCTTTTGAG CTGTATGCAT TTGGCTATTA 300
 TGAATAAAGC TGCTTTGAAC ATTCCTAAA 329

SEQ ID NO:7281

SEQUENCE LENGTH:270

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08520

SEQUENCE DESCRIPTION:

GATCAGAGAA GACCCCTGAG ATGAATCAGG GCCCTGCCCT GNCCACCAAC TCTTTTCTAG 60
 CAATCATGTC CTGCCCAGGC TGCAGGCTTC ATTAGGCTCC TGCTGTCTC CATGATGTGA 120
 CTAGCACAAG GTGTATATAT GTTTGTACC TCTGCCGATG GCTGTACATA GTGTATGAAA 180
 GTTATTTAAG CCCCATGTTG TACATTTCTG TTCCTAGATT GGATGTGTGT GTTCTAAGAA 240
 GTTGTCATAA ATAAACCTG AATGACCAAA 270

SEQ ID NO:7282

SEQUENCE LENGTH:195

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08521

SEQUENCE DESCRIPTION:

GATCCAATTA TTAGAAAAGC ACTCTTAAGA GTGCAATGAA TGTGATTGA AAACCTGAAA 60
 GCAGCCTAGA TTTTACTATA TTTGTTATTG TTATCAATGG GGGTTTGCA ACCTTAAAAA 120
 TTATTTTNA TGTAAGATA CCTAGGAATT TCTAATAACT GTATTGTTT TAATAAACTT 180
 TGACTTTGCT GTAAA 195

SEQ ID NO:7283

SEQUENCE LENGTH:386

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08522

SEQUENCE DESCRIPTION:

GATCTACCTC AGTTAAACAG TTGGGTGCTA TTAATAAGTC TGTCAAATTA AATTGGAAAA 60

EP 0 679 716 A1

5 AGTAACCAAA CGGTGAGATA CAACTCCACA TGAAACTGA AATTGTAATT TCCGTTTATT 120
 TAATGATATT TTNATTTATT TGTGCCTTTT ATGTTGAACC CCAATGCATT GAAAAANTTC 180
 AGTATGAAAC AGTACATATT TNATTTATAT TACAGGTGGG AGAAAAGTCC AATTGGTCAT 240
 GGAATTTGAT AGACTTTTCC CCAGCCAAC TGTACAGTGT ATTATAATCC CGACTGCCCC 300
 CCTGTGAAAA GANAAAAAAA ATTGTCCANG GGCAATTTN CCATTTCNA ACCANTANNG 360
 NCCANNCTCC TTAAAAANGN GNGNNN 386

10 SEQ ID NO:7284
 SEQUENCE LENGTH:187
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08523
 15 SEQUENCE DESCRIPTION:
 GATCAGTTAG TAGGCTTTCG NTGTCTTCTC TTTCAATACA TGTACATCTT TACTGTTTGA 60
 AAAGTGTTAC AGCTGTCAA GAATCTTCAT GGACCTGAAG ATAATTINTT GTGAAGTTGA 120
 ATGCAAGNT ACTGTCATTC ATAGTGTTAA TATCAAANTA CCAGGAATNT TNACTTTTNC 180
 NACCTNN 187

20 SEQ ID NO:7285
 SEQUENCE LENGTH:113
 SEQUENCE TYPE:nucleic acid
 25 TOPOLOGY:linear
 CLONE:HUMGS08524
 SEQUENCE DESCRIPTION:
 GATCTCAGAA ATTGTATCTN AGTTGGTATC AACCAAATGG AGTGACTTAG TGTACATGAA 60
 AGCGTAAAGA GGATGNTGG CATTNNACT TTTGGCTTGT AAAGTACNGG CTN 113

30 SEQ ID NO:7286
 SEQUENCE LENGTH:120
 SEQUENCE TYPE:nucleic acid
 35 TOPOLOGY:linear
 CLONE:HUMGS08525
 SEQUENCE DESCRIPTION:
 GATCAAAGAG TAAGCCACAC ACGGATAATC AGTTACTAGG GATGGAGGTG TGAGGGTTCA 60
 TTATATTATT CATTTTACTG TTGTATATGT TTGAAAATNT CTATAATAA AAGCTTTAAA 120

40 SEQ ID NO:7287
 SEQUENCE LENGTH:65
 SEQUENCE TYPE:nucleic acid
 45 TOPOLOGY:linear
 CLONE:HUMGS08526
 SEQUENCE DESCRIPTION:
 GATCATGTGT ATGAATTGTT GGTGCTCTAA AGAACAGCAC AAATAAAATN ATTTTCAAAT 60
 TTAAA 65

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EP 0 679 716 A1

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SEQ ID NO:7288
SEQUENCE LENGTH:110
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08527
SEQUENCE DESCRIPTION:
GATCCGNC CG CCTCAGCCTC CCAAAGTGCT GGGATTACAG GCATGANCCA CTGCACCCGG 60
CCCCATTCTC CACTTTAGCC TCAGGCAGAG AAGACNGCGC GATGGGGNNN 110

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SEQ ID NO:7289
SEQUENCE LENGTH:296
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08528
SEQUENCE DESCRIPTION:
GATCTGCGAC ACTGTGGACT GAACACACTG AAGCTCTGAT GGGAAAACCT GGTGACTGAT 60
ATAGTTGTTT AGCAATAATT CATAGTCTGT GATGAAGAGT AGTGAATACA ACACACAACC 120
AGGCAGCCTT CTGACTATA CTTTGACAT GNTTCTTTA GGAATTCAT CACACATTAT 180
AACCAGTTAG TGCCTTCTAG AAGAATGGCT TTCCTTTNTN CTACACAAAN TTTGAATTAT 240
ACAAGNNTCT AAATATAATA CCTTNAATA AAAAGGNTAA TTNNGNCCCT CTGAAA 296

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SEQ ID NO:7290
SEQUENCE LENGTH:381
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08529
SEQUENCE DESCRIPTION:
GATCAGGTCC AAGCAAAACG TCCAGAGTGC TACAAAATGT TCGTTCTCA GTCCAAAAG 60
AAGTGGAAAA GAATCTGAAG TCATGCTTGG ACAATGTTAA TGTGTGTCC GTAGACACTG 120
CCAGAACACT ATTCAACCA GTGATGGAAA AGGAGTTTGA AGACGGCATC ATTAAGTGG 180
GAAGAATTGT AACCATATTT GCATTGANG GTATTCTCAT CAAGAACTT CTACGACAGC 240
AAATTGCCCC GGATGTGGAT ACCTATANGG NGATTTCATA TTTNTTGGC GAGTTCATAA 300
TGAATAACAC AGGAGGATGG GNTAAGGCAA AACGGAGGCT GGGTATGTGT GATGGGAAAA 360
CTTTCTTCAT TGGTNCCTTC N 381

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SEQ ID NO:7291
SEQUENCE LENGTH:51
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08530
SEQUENCE DESCRIPTION:
GATCACCCCC GCCGACGGCC CCGGGCCCCG ACGGCCCGGA AGTTCGCAA A 51

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SEQ ID NO:7292
SEQUENCE LENGTH:347

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08531

SEQUENCE DESCRIPTION:

GATCATTCTG TTCTTTCAAG GAGAAATAAG CCTAAAAGAA GAAAAACAAA AAAAATTCTG 60
TATAAACTG TAATCCTTTG TATTCATGTT TACAGTGCTA TTACTATAAT NCAAAATTAT 120
GTATGTGACT TAGAGTTATA TAATCATAAT TTATGTTTAT TTCAAATATC TAAGTTTATT 180
GCTTGGATTT CTAGTGAGAG CTGTTGAATT TGGTGATGTC AAATGTTTCT AGGGTTTTTT 240
TAGTTTGT TTATTGAAAA NTTAATNAT TTATGCTATA GGTGATATTC TCTTTGAATA 300
AACCNATAAT AGAAAATAGC AGNCACCATA ANCATCTTTG TAAATNN 347

SEQ ID NO:7293

SEQUENCE LENGTH:122

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08532

SEQUENCE DESCRIPTION:

GATCCAGAAA CTTTCCCAA GAAGCTTGA CATGGATTGC TGGTAAACAC AAAAAGTGA 60
TGGAGGGCAG AGCAGCCGGT TGACTCCCC TTGTCAATAT CTAAAGTGG AAAAGCTTGA 120
AA 122

SEQ ID NO:7294

SEQUENCE LENGTH:138

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08533

SEQUENCE DESCRIPTION:

GATCATCTTT CCTGTTCCAG AGAAGTGGGC TGGATGTCTC CATCTCTGTC TCAACTTTAC 60
GTGTAAGTCTG CTGCAACTTC TTAAGTTCCC TAACTNAAAG ANCGACTCTG ACAGAACTN 120
TNTTNCNCA ACAGTTTN 138

SEQ ID NO:7295

SEQUENCE LENGTH:211

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08534

SEQUENCE DESCRIPTION:

GATCCTTCCG GNGGAGTGTG GGAGAAGAGA GCTTGCCGAC GATGCCTTCC TGTGCAGAGC 60
TTGGGCATNT CCTTACGCC AGGGTGAGGA AGACACCAGG ACAATGACAG CATCGGGTGT 120
TGTTCTCATC ACAGCGCTC AGTTAGAGGA TGTTCTCTT GGTGACCTCA TGTAATTAGC 180
TCATTCAATA AAGCACTTC TTTATTTTAA A 211

SEQ ID NO:7296

SEQUENCE LENGTH:152

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08535

SEQUENCE DESCRIPTION:

5 GATCAACGGT ACTTGTTTAA AAGAAGCTTA TGTGGACTTT GTCTTTTCTC GTTGCCACAA 60
CCTCTTTTAA TGTGCCTATA GTCCACTATA GCATGCATAT CTCAAATTGC AATTCCTTGC 120
AATTCTGGAA TAAACGCTTT ATTTGGAGA AA 152

SEQ ID NO:7297

10 SEQUENCE LENGTH:381

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08536

15 SEQUENCE DESCRIPTION:

GATCTGCTGA CTTATATAAA GCTTTTTGAT TCCTACTAAG CTTTAAGATT TAAAAAATGT 60
TCAATGTTGA AATTCTGTG GGGCTCTATT TTTNCTTTGG CTTTCTGGTG AGAGAGTGAG 120
GAAGCATTCT TTCCTTCACT AAGTTTGNT TTCTTGCTTT CTGGATAGAT TGATTTTAAG 180
AGACTAAGGG AATTACAAA CTAAAGATTT TAGTCATCTG GTGGAAAAGG AGACTTTAAG 240
20 ATTGTTTAGG GCTGGGCGGG GTGACTCACA TCTGTAATCC CAGCACTTTG GGAGGCCAAG 300
GCAGGCAGAC CACTTGANGG AGTTCGGGNC CAGCGTGGCC ACCGTGGTGA ANCCCTGTNT 360
CTNCTAAAAA TNCAAAATTT N 381

SEQ ID NO:7298

25 SEQUENCE LENGTH:379

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08537

30 SEQUENCE DESCRIPTION:

GATCAACAGC CTCCTCTCTT GGGGACTCTC AAGAGCCTGT TTTTCATCTAG AAGTAGTAGT 60
TTGATTCTGG TTCCCTCCT ACAGTGTGTC CTCCTCTCTT TTGCAGCTCC GTCATTACCA 120
TAGGGGACTT GGTTTTAGAC TCTGATGAGG AAGAAAATGG CCAGGGGGAA GGAAAGGTGA 180
35 GTGGGAAGGA GCAGAAAGCT GGGAAAGGGG ATGGGTAGAA CAAGACTGAG AAATCCACAT 240
GCTTCAGANT TCAGAGGGTT CAGGGAATNG TTTCCGATAG TAGGCTCTCC CTGCTCCCTT 300
CTCTACAGGA ATCTNTGGAA AACTATCAGA AGACAAAGTT TGACACCTTG ATACCCACTC 360
TNTGTGAATA CCCTACCCN 379

SEQ ID NO:7299

40 SEQUENCE LENGTH:82

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08538

45 SEQUENCE DESCRIPTION:

GATCATTACA AACAAAACT GTAATTTTNT TATATTTGAT TCAATGGAAT TTACCTAAAA 60
AATAAAGACT AAAAATGTGA AA 82

50 SEQ ID NO:7300

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EP 0 679 716 A1

SEQUENCE LENGTH:225

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08539

SEQUENCE DESCRIPTION:

GATCCCTTTA TTGACAGCCA GTTTGAGGAC ACTGTCCCAG CATCTCTAAT GGAGCCTGAG 60
CCGGTGTGAG GACCAGGATG TCTTTTCCCA GCCCAAGAG ACCTGTGCT GCTTTCTTGT 120
AATNATGGGG CTCCCAGAG TCTGCGTAAC AGTCTCCAC TGGCTGGCTC ACCCACAGGT 180
GCCATGTGCA CACTCCTGGT TTTCAAACAA TTCTCTGGAT TTAAA 225

SEQ ID NO:7301

SEQUENCE LENGTH:216

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08540

SEQUENCE DESCRIPTION:

GATCATGCAT GCCAATCCCT GTTGCCGCAT GGAGCTCCTC AGCCCACTGA CCTCTCCGTG 60
CCTGGTGCAG GCCAGGCCCC CGTCTTCCGC CTGCCTCTGC TTCCCCGTCA TGCATGGTGG 120
TGGTGTCTTCT ACGGTGTCTG GTTCTGTGCC CGTCTCTGAG ACAGTCTCTG TGTGGAATNT 180
GCCTTAAACT GAAGTAAATN TGGTTCTTTT AGTAAA 216

SEQ ID NO:7302

SEQUENCE LENGTH:374

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08541

SEQUENCE DESCRIPTION:

GATCAATTCA TGCAAGATAT AATCATATTT TTAAATTAAG AAAATTTTTT TTGTAGAGAT 60
GGGGTTATGC TGTGTTGCCT GGCTGGTCTC AAACCTCTGGG TCTCAAAGCG GTCCTCCTGC 120
TTTGGCCTCC TAAGGTGCTG GGATTATAGG CATGAGCCAC TGTGCCAGGC CTGAGGATAT 180
TATCATTATA AATATATATG TACCCACAC TGGATGACCC AGATATAAAG CAAATATTAT 240
TAGATTTAAA GGGAGAAATA GACTCTAATA CAATAGTAGT TGAGGACTTC ACTCTACTCT 300
CAGCATTGGA CAGATTTTCT AGACATAAAA CCAATAAATA TTGGATTAA ATTGCACTTT 360
ACACCAAATG GTCN 374

SEQ ID NO:7303

SEQUENCE LENGTH:113

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08542

SEQUENCE DESCRIPTION:

GATCCCATTT CTATGAAATG TCCAGGGCAG GCCCAGCCAC AGAGACAGGG AGTAACGTGG 60
TGGTTTATTA TAAATTTTTC ATGAAAATAA AATATACAGT ACTCTCGTGN AAA 113

SEQ ID NO:7304

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08543

SEQUENCE DESCRIPTION:

GATCTGCACA GCCTCTAGAG GCCTCCCAGC AAATGCGGGG AGCCATGCCC CCAGGGTCTA 60
 CACACTCTCG TTCATCAACA TCACAAC TGG AATTCGGGAT TTGTGAAGTT TAGAGCTGNA 120
 CAGACTGTGA CAGATTATGA GTCAACACGT ATATTTTCTC TTTCAAAATA ATAATATTTT 180
 GTTTTTGACT TTTTACTAAG TGAATATTAT TTTTAAATC TGCCTATATA TTGGAACCTC 240
 TATTTTATAA TAATAATGAT AATAAATCAG TACCCAGAAG TATAAA 286

SEQ ID NO:7305

SEQUENCE LENGTH:116

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08544

SEQUENCE DESCRIPTION:

GATCTAAATN ATGTACTTGT TAGTGTATTC ATTCATATTG ATTGTAAAGG ATTATTTTTC 60
 ACTCAGTACT GATGTCCTTG GAAATCTTAC CTGGAAACAT GTTTGCAAAA AACAAA 116

SEQ ID NO:7306

SEQUENCE LENGTH:137

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08545

SEQUENCE DESCRIPTION:

GATCCCAGT CTACCCTGGA AATNCCCAGG CACAGCATGC CTGGAGGTGT GAGCCGCCTC 60
 TTAGAAGCTT TATAATCCCG GATAATNCTG CTGTAATTGG CATTGTTTTT CCATTACAAT 120
 CCTTTCATTA CCTGAAA 137

SEQ ID NO:7307

SEQUENCE LENGTH:317

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08546

SEQUENCE DESCRIPTION:

GATCCCAGCT TGGTGGGAAA GTGCAGAAGA ATTGCAAAAC TGACATCCCA TNTNACAGCA 60
 ATAGTGACCT TTATTTAAAT TGTGTGTGA TAGTTTATGC TTCTTAAATC ATTTTCAAC 120
 NTAACAGCC AATTTCTAAG CAGACAGGAA AACTAAATAA TAAGTTNATT NATATAACAA 180
 AGATGCAGGT TCCTGCTCAT TCCAGTAATG TCTTTGAAAG CANNNCTNAT ATTTNTTTTC 240
 TAGATTATCC CTGTGAATAA TTGAGANCTT TTTGGAGTCA AGNATGAATA AAGGTGTGGC 300
 AAGGNTATAA ACANNAN 317

SEQ ID NO:7308

SEQUENCE LENGTH:337

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08547

SEQUENCE DESCRIPTION:

GATCTGTCAG AAAAGACATA TTTTGTAGAT AAAGGAGTTT CTAAATAATG AATTATTGTT 60
TAGTACCAGC TTTTGAATC CTTACATTTT AAGGAACTTG AATAATTTAA ATATTAGATA 120
ATGTAACGCC TCATAAAAC AGTGTCTAGA CAAGGGCCAA TATTGCATTG AGATTGGATG 180
AACATTTTAT TTTGCATATA ATTGTTGTCT ATAAACAAA ATACTAATAA TAATGGCTAC 240
CNGTTAGNCA NTTTAGNNCN TGTCTTACTA TTTACCNNTG TGGNAAAACA TTGCATGNTT 300
TAATNTTTTT CTAAATAAAT ATTNTNCCAT TANCNNN 337

SEQ ID NO:7309

SEQUENCE LENGTH:389

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08548

SEQUENCE DESCRIPTION:

GATCCCACCA ATCCTCGTTC CCCCTAAACA TGCTAACAAA AATCCCTTAT TGTGGGTATT 60
AAAAATAATA CAGTTACACT GTATGCATAT TTATGTGCTC CTTTGTCTG GTTTNCTTT 120
TCATCATGTA TAAGCTGAAT TCAGCATTAG TTTCTCACAT CTTCCCCCAG GTATCCCCAA 180
CAGAATTTT ATGTCCCAGC TTGTATTAAA TAGAAGTGAA ATATTAAGGA AAATAAGGAA 240
CTTGTGCAAC TTTTNNATG CATTGTTCTC AACCATTTAA TTTATTGAAA GGAGATGCTG 300
CAACAGTTCT TGATTTAGCA GCAGTTATTC TCTTGTTTAC ATAGTTATGG TTTTNTGTTG 360
TTGGTTTTTG CTCTGGTACT GGGAACCGN 389

SEQ ID NO:7310

SEQUENCE LENGTH:367

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08549

SEQUENCE DESCRIPTION:

GATCCAGGTC TTCTGATTCT TATCCAGTG TCCTTTCTAG CATACCATGT TGCTCTAAA 60
GATTGCAGCT CCTTATTAC TAGAAAATTG TTCCTGCCCA ATCTACATCT CCACCTCACC 120
CCATCTTTTC TTAAGCACTA TGTTGTGTT TTTATCAGTA TTATATTCAT TGTCTTTGGA 180
ATACATGTTT TTGTTTGTGT TTGGAAAAA AATCTCTTTT ACCAGCTTGC ACTCGGACCA 240
ACTTGAAAAA AAAAANGCTT AAATGTTTNN GCTATGTNCA GTTTAAAAAT GTGAAGTTTG 300
TNGCTTTANC TTTTGTAGG AAAATCTANT ANCACTGGCT TAAGTGCTGA CTTGAAATGC 360
TATCCAN 367

SEQ ID NO:7311

SEQUENCE LENGTH:56

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08550

SEQUENCE DESCRIPTION:

GATCCAGCGA ACCACTGCAC TCTAGCTGGG GAGACAGAGC AAGACTCCGT CTCAAA 56

SEQ ID NO:7312

SEQUENCE LENGTH:92

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08551

SEQUENCE DESCRIPTION:

GATCTCTTCC CTTAGGAAAT ATCAACATCT TTATGTACAT GTCAGAGTCA TGGTGTGATG 60
ACGTACTTTG AAGCATGGAG TCAGTTTCA AA 92

SEQ ID NO:7313

SEQUENCE LENGTH:368

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08552

SEQUENCE DESCRIPTION:

GATCCACTTA TGAAGTGAAC ATTCTCCATG AACAGGTGGT TGGATTGGTA TCTGTCATTG 60
TAGGGATAGA TAATAAGCTC TTCTTATTCA TGTGTAAGGG AGGTCCATAG AATTTAGGTG 120
GTCTGTCAAC TATTCTACTT ATGAGAGAAT TGGTCTGTAC ATTGACTGAT TCACTTTTTC 180
ATAAAGTGAG CATTATTGA GCATTTTNC ATGTGCCAGA GCCTGTACTG GAGGCCCCCA 240
TTGTGCACAC ATGGAGAGAA CATGAGTCTC TCTTAATTTT TATCTGGTTG CTAAAGAATT 300
ATTACCAAT AAAATTATAT GATGTGGA AAAA AATTGCGGGC TTGNANGNTN 360
GGGGTCAN 368

SEQ ID NO:7314

SEQUENCE LENGTH:297

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08553

SEQUENCE DESCRIPTION:

GATCACACCA CTGTACTCCA GCCTGGATGA CAGAGTGGAG ACTCTGTTTC AAAAAACAG 60
AAAAGAAAAT ATAGTTTGAT TCTTCATTTT TTAAATTTG CAAATCTCAG GATAAAGTTT 120
GCTAAGTAAA TTAGTAATGT ACTATAGATA TAACTGTACA AAANTTGTTT AACCTAAAC 180
AATCTGTAAT TGCTTATTGT TTTATTGTAT ACTCTTTGTC TTTTAAAGAC CCCTAATAGC 240
CTTTTGTAAC TTGATGGCTT AAAATTCCTN ATTAATNCTG CCATTTCAN TTCCAAA 297

SEQ ID NO:7315

SEQUENCE LENGTH:86

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08554

SEQUENCE DESCRIPTION:

GATCTTTGAA TGAGCTTTT AAGGAAGAAA TATTATATAT TGTGTTTAA AGTTTATTGA 60
AATAAAGAAT CATTTAAATC TTCAA 86

EP 0 679 716 A1

SEQ ID NO:7316
 SEQUENCE LENGTH:356
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08555
 SEQUENCE DESCRIPTION:
 GATCCAAATT GTGAAGCCGT TGGTCAGGTG CCTGGCTACA CACAAAATGG TGGAGGCTTA 60
 GCCTAAACAG TTTCTTCATT TCACATTAAA ATCCTAATTN NTGGCTCCTC TCAAAATGAG 120
 TTCTAGAAGC ACNCGNTGTA GATTCCCATG TGGACACAGC CTCAGGAGG TGAGCAGCAG 180
 CTCCCTGCAG CCAGCTGGAG CCTGCCAGTT TTCCCTAGTG CCCAGCACTC CTGAGCTTCT 240
 GCCCTCCGAG GGATTCCCGA AGTTTTTCT AATGGGNTTA GTGGNGNCNC CTTGGGGTT 300
 GCTTNGGTTT CNTTTNGTTA NGNGCCCTTG NTTTTAAAT AAANCNGGGC CATTNN 356

SEQ ID NO:7317
 SEQUENCE LENGTH:349
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08556
 SEQUENCE DESCRIPTION:
 GATCTGTGGG ACTGTCTGGG CCTGTACTC ATCCTGCTAT CAATTTCTTA TTAATTAATC 60
 TTGATGATTC TNATTAATTA ATCACATTG CAGGAAATTC AGATGAGGCA AGAAAATTTN 120
 ATTGGCCTGG GNNNGACTGA AAGCATTCCA AATTAGGCTT AGACTGTGCA AAGGGCTTAG 180
 CTAAGTTATC GAGCTTAAAA CCCGTCAATT AANCAACAT TATTGANCA GTTACTGCAT 240
 GCCACGCACT GTGTTGGGCT TAGTAATAAA ANNNAGNAAA GGTAAGNGCT TGTNCTNGCC 300
 TAAATTAAG GGTCCCCGGG GGTTTTTTCT NGAGGGGGCN NCTGCCCN 349

SEQ ID NO:7318
 SEQUENCE LENGTH:351
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08557
 SEQUENCE DESCRIPTION:
 GATCACCTGA GGTGAGGAGT TCAAGACCAG CCTGGCCAAC ATGGTGAAAC GCTGTCTCTG 60
 CTAANTAC AAAANTTAGC TGGGCATGGT GGTGGGCACC TGTAATCCCA GCTACTCGGG 120
 AGGCTGAGGC AGGAGANTCG CTTGATTCCA GGAGGCAGAG GTTGCACTGA GCCAAGACTG 180
 CACTGTTGCA CTCCAACCTG GGCAACAAGA GCAAACTCC GTCTCAAAA AAAANGTGT 240
 TAATTATTCT AAAAAAATT CAGCCAGGTT ATTCCAAATT NTAATTGAAA NCNGGGGGCC 300
 CAGGTTAANC GGGGCCTTAC CCTTCCCTTN GCNCTTTGGA NTTNNCCAT N 351

SEQ ID NO:7319
 SEQUENCE LENGTH:333
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08558

SEQUENCE DESCRIPTION:

GATCAGCAGT GTTGACCATT TATGCTGCAT AGCTGGTATT ATAGCCTTAT TAGTTGTGTG 60
 GTTGACCCTT GGGGTATACA AATGTCAGTC TGAGTGGTGN CTTACTCCTT TGTTTATAAG 120
 TGAATGATTG TGCATGTTTT GTATGTCATA GTATGTCGTC ACATAAAAGG GAGGGAGCGA 180
 AAAACCATTA CATTAAGATA ATATTGGACC AACTACTTA CTTGCTCTAA ACAGTTACTT 240
 GTACCCCTTA ACCTGTCTTC AAAAGTTGCA TATAGTTACA GTAGTGTATA AATTAAATAT 300
 TGTGGAAC CANNAANN NNNNANNNN NNN 333

SEQ ID NO:7320

SEQUENCE LENGTH:192

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08559

SEQUENCE DESCRIPTION:

GATCAGCTGA AGGTTTCATGG GTTTAAGTG CTTGTGGCTC ACTGAAGCTT AAGTGAGGAT 60
 TTCCTTGCAA TGAGTAGAAT TTCCTTCTC TCCCTTGTC CAGGTNTAAA ACCTCACAGT 120
 TGATAATGTA CCATTGGGNN CCGTTTAAAC TGGGCCTAGG NACCCCTCA NGAATAACCT 180
 GNAAGGGCCA AA 192

SEQ ID NO:7321

SEQUENCE LENGTH:55

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08560

SEQUENCE DESCRIPTION:

GATCGTGCCA TTCCACTCCA GCTTGGACAA CAAGAGCAA ACTCCGTTTA TAAA 55

SEQ ID NO:7322

SEQUENCE LENGTH:78

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08561

SEQUENCE DESCRIPTION:

GATCAGACCC CCACAATATG TCTCAACCTC TGTGAAGGG CTAGTGATGT TAAAAAAC 60
 CTGTTATTTT CAATCAA 78

SEQ ID NO:7323

SEQUENCE LENGTH:375

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08563

SEQUENCE DESCRIPTION:

GATCTGGTAT TTCCTGGACT AAATCCCCT TGGGAAGAC GAAGGGATAN TGCAGTTCCA 60
 AAAGAGAAGG ACTCTCCAG AGTCATCTAC CTGAGTCCCA AAGCTCCCTG TCCTGAAAGC 120
 CACAGACAAT ATGGTCCCA ATAAGTACT GCACCTTCTG TGCCTCAGCC GTTCTTGACA 180

EP 0 679 716 A1

5 TCAAGAATCT TCTGTTCCAC ATCCACACAG CCAATACAAT TAGTCAAACC ACTGTTATTA 240
ACAGATGTAG CAACATGAGA AACGCTTATG TTACAGGTTA CATGAGAGCA ATCATGTAAG 300
TCTATATGAC TTCAGAAATG TTAATAATAGA CTAACCTCTA ACAACAAATT AAAAGTGATT 360
GTTTCAAGGT GAAAA 375

10 SEQ ID NO:7324
SEQUENCE LENGTH:271
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08564
SEQUENCE DESCRIPTION:
15 GATCCACCTG CCTCGGCCTC CCAAAGTGCT GCGATTACAG GCATGAGCCA CCGCGCCTGG 60
CCGCTTTCGA TATTTTCTAA ACTTTAATTC AAAAGCACTT TGTGCTGTGT TCTATATAAA 120
AAACATAATA AAAATTGAAA TGAAAGAATA ATTGTTATTA TAAAAGTACT AGCTTACTTT 180
TGTATGGATT CAGAATATAC TAAATTAACT TTTTAAACA CAACTTTTAA AAAATGTATC 240
AAAATAATAA ACGTGTTCTG ATATTTTAA A 271

20 SEQ ID NO:7325
SEQUENCE LENGTH:102
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS08565
SEQUENCE DESCRIPTION:
GATCCCTCGA GTTCAAGAGT TTGAGACCAG CCTAGGCAAC CTTGTCTCTA CAGAAAAATA 60
AATAAATAAA TAAATAAACA AACAAACAAA CCAAAAATTA AA 102

30 SEQ ID NO:7326
SEQUENCE LENGTH:216
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
35 CLONE:HUMGS08566
SEQUENCE DESCRIPTION:
GATCATGAGG TCAGGAGACC GAGCACCATC CTGGCTGACA CGGTGAGACC NCGTCTNTAC 60
TAAAAATACA AAAGCAAAAA TTAAGTGGGT GTGGTGGCGG GTGCCTGCAG TCCAGCTAC 120
TNGGGAGGCT GAGGCAGGAG AATGGCATGA ACCTGGGAGG CGGANTTGCA TTGAGCTGAG 180
40 ATGGCACCCAC TGCACTCCAG CCTGAGCGAN NNNNNN 216

45 SEQ ID NO:7327
SEQUENCE LENGTH:176
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08567
SEQUENCE DESCRIPTION:
50 GATCTGCTCT CTGTTGAGAG TTGGTAATCA TTGGTTTGAA ATGTGATGAA ACCACTCAAG 60
CCAATGAAGG TGGGTGTGTA GGTGGGGAGT ACTTTGCCAT AATATTTTAA AACATTACCT 120

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GGTTAGAGTT CTAAGTGGTA CTTATTTTGG TTTGGTTAGG GGAAAGCCTG AATAAA 176

SEQ ID NO:7328

SEQUENCE LENGTH:106

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08568

SEQUENCE DESCRIPTION:

GATCTCTGGC TGTTTGATT TTTTATATT GTATTTTAA AATNTGTTAA ACAGTGCCCT 60
GTGAGCACCA AGTACCACTA GATGAATAAA ACGTATTATA TCTAAA 106

SEQ ID NO:7329

SEQUENCE LENGTH:379

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08569

SEQUENCE DESCRIPTION:

GATCCGGCAG GCAGAGAGGA AAAAGGGAAC TTGTACCAAG CACAAACAGG NTCTAGGCAT 60
AGTGCTGTGT ACTTAGGTTT TTCCTGTGCT TCAGTCTTTC CAATAAATCC AGGGGTAGG 120
TTTTAATCTN NCTATTTNAT AGTTACAGAC CATTTTAAAT GGACAAGCAG CCCATTAAAA 180
GATAAAAGTA CTTAGATGGA CTAGCAGCTT CTCTACCAA TTATTCACC ACACAAAGNT 240
AACATCATTG TGAATTTTGG TTTAGTTAAC ATGCATTCCC TGGCCTGCCT CACACCGGAC 300
CTNTACCCGG TACCATGCAA ATGCACATCT GCTGCTACAC TTGTGGGATG CTGGTTTAGG 360
TAATTNTTTT TTATAATNN 379

SEQ ID NO:7330

SEQUENCE LENGTH:271

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08570

SEQUENCE DESCRIPTION:

GATCCCAGCA GAATACCAA ATCCTATTTT TTTGACTGA GTATTTGTAG ATGCTTAATG 60
ACTGAAATNA ATTTGGAGGC ACTGATGAAA GTAATTTTTT NAAAGTTCTC AGGTACTGTT 120
CAATTATTTA ATGTTAAGTT TAGTATCAAG ATACAGTTGT TTNAAAATG CCAAAATGCT 180
GTTTATNATA CAGAATATTT NATTACATTT GCAATATCTT TGTATATAGT GATTTTNTNC 240
TTGATAATAA ATGGAAAANT TCTAAACCAA A 271

SEQ ID NO:7331

SEQUENCE LENGTH:382

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08571

SEQUENCE DESCRIPTION:

GATCGACCTG CCTCTGCTTC CCAAAGTGCT GGGATTACAG GCGTGAACCTA CCGCGCCTGG 60
CCCTCAGATA TTTTAAATT TGAATAATTA GAGTATATAT TAATATCAAG TNGCAGAAAC 120

EP 0 679 716 A1

5 AGAATGAAGT GTTATAAAGG CGTTAAGTAG TAGGGNGGCA TAAAACTTAA ATCTTTAAGC 180
TAGAAATAGC TTGAGGAATT ATCTGATGCA GTACTCTGTT GTAATATAAC AAAAATGGAA 240
TCACAGTTTT ACAAATCAGG CAGCTGCAGC TCAGNGAGGT TAAATTTCCA CAGACCTGTT 300
TATGCAGCGC GGTAANTGAG GAAGTAGACC TAAATTTGG GGCCTCTGCT ACTCTGGGCC 360
ATTACCTCAG CCTCGCTGGC AN 382

10 SEQ ID NO:7332
SEQUENCE LENGTH:315
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08572
SEQUENCE DESCRIPTION:
15 GATCTCACAG GAATCCTTCA ACTGCTTTCT ACCAAGCGTT CCATTTGAAC ACGTTAAAGG 60
AATCAAAGAG CCTCTGGGAT AGGGGGGAAA NCACATGGTA TCTAGCATGG GACAAAGCAG 120
CAAAATAAAC CAAAAGGACA ATCTCTATTT TATATAGAAA ATAAACATAC TTTTCTGCAT 180
CCTGTGTACT TTAAGTATAT CTATATTTTT GAAGGGTTCA TACTGTGTTG AATTTTTCTT 240
ATGAAATAGT CACTTCCCCA GTGTATTTA ATGCAATGC ATATTCTATA AAATAACCTA 300
20 TATTTTAATT TAAA 315

25 SEQ ID NO:7333
SEQUENCE LENGTH:104
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08573
SEQUENCE DESCRIPTION:
30 GATCTTTAAT ATCTGATATA TTCCTGGTAC TCGTACTGAT AAGGGATTAT TGGAAGTCAG 60
TCACAGAATT TGGAAATAAA TTCTAGTCTN TCCTTAGCTN NNNN 104

35 SEQ ID NO:7334
SEQUENCE LENGTH:140
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08574
SEQUENCE DESCRIPTION:
40 GATCTACCTG AGGCTCAGTC CTCATCTGTA AAATGCTGAT AAAAGGACCT ACCACCTAGC 60
GTTGCTGTGA GGTGTCAGTG AAATAAAATG TAAATGCTT AGCACAGTGA ACTTTAATAA 120
ATGGTAGTTG TGGTTGCAAA 140

45 SEQ ID NO:7335
SEQUENCE LENGTH:93
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08575
SEQUENCE DESCRIPTION:
50 GATCTAAAGT CTAATAAACA AATCCTTAGG TTAAGAAGAA AATAATGCAA TAAATAAAAT 60

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GTTATAAATA AATGACATAT TGCNACTTTT AAA

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SEQ ID NO:7336

SEQUENCE LENGTH:96

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08576

SEQUENCE DESCRIPTION:

GATCATACCC ATGAAACCCA CTGTTTAGAT TTAAAAAGT TCAATGTTTT TCTGTATTN 60

ATCTCTTCAT TTTAAAAATA AGTTTGCAC ATGAAA 96

SEQ ID NO:7337

SEQUENCE LENGTH:224

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08577

SEQUENCE DESCRIPTION:

GATCTAAAAG CAGATGCTGT AGCTGCCTAC TTTAACCCAT TTTCAACCT GTTTGTTTTT 60

TAAAGGGCTT CACTAAGGGT NNNNNATGTA CCATTGTAGG GGGCAATTTT AAGTCAGCTA 120

AGGCAATAAC CTTATGCATG AACATTTCCC AGACTTTCAT GAAGCTGTG AGGTCCTAGG 180

CAATTAATGC GGCAGTTGTG ATAAATAAAA ACATCTCACC TAAA 224

SEQ ID NO:7338

SEQUENCE LENGTH:175

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08578

SEQUENCE DESCRIPTION:

GATCNATATT TAGTGACTGC AACATGTTTA TACCACTGAT TCAAATTCCA NNNNTGATGA 60

AGTTATACAA ATAATGCATA TATTGATAAC TTTTATTGCA AAAATGTAAA TTTAAAACTT 120

GTATAATGTT CTTGTGCTTT TTTAAATAAA ATATATGTGT ATATTTAAAA AGAAA 175

SEQ ID NO:7339

SEQUENCE LENGTH:143

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08579

SEQUENCE DESCRIPTION:

GATCTGAGGA TGGTTATAAA TACTGTAAAT ATTGTAATGT TATGAATGCA ANNNTATTTG 60

AAAGCTGTTT ATNATNATAT CATTCTGAT AATGCTATGT GAGTGTITTT AATAAAATTN 120

ATATTTATTT AATGCACTCT AAA 143

SEQ ID NO:7340

SEQUENCE LENGTH:106

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08580

SEQUENCE DESCRIPTION:

GATCACTTGA TTTTAAAGAA ATNAAAAATT GCACTAGGTA GAAATAGTGT ACCAGGCAAT 60
AGTCCAGCTT TAGTTTNTG TCATTTTNG TCTCTATTTA AAGNNN 106

SEQ ID NO:7341

SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08581

SEQUENCE DESCRIPTION:

GATCTCAAAG CAAATCTGTA TTTTAATCTC TCAGGTGTTC ACTAGTATTA CCAGTGACTA 60
TGGGAAATAA TAAAGGAAA TCTTGTCAT AAA 93

SEQ ID NO:7342

SEQUENCE LENGTH:193

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08582

SEQUENCE DESCRIPTION:

GATCTTCGCT CCAGCTTTCT GCATCTGAGC TGGAAAAAGA GCAAAAACGG GTAAAAATTG 60
AATCCAAATA TTGTCGNATG TATGCCTAAT TTAGAATGAA ATCTTGGGGT GGCAGACTCT 120
GGAGTTCTCT GTGAGGTGTG GCATTCTTTT NTCTAAATCA CTTATTGGCC AATGTATGTG 180
TCACTTTGTT AAA 193

SEQ ID NO:7343

SEQUENCE LENGTH:137

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08583

SEQUENCE DESCRIPTION:

GATCTATAAT AAGAGCTCAA TAACTNNGTC AAGGAAAGCT CTAATATATG CAGTGATGGT 60
TTATGAAAGG GTGTGGCAAT NTNAAATNTA TATTGTGTGT GATGTTCAAA TAAAGTGGA 120
TCTACATNCA TGTGAAA 137

SEQ ID NO:7344

SEQUENCE LENGTH:389

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08584

SEQUENCE DESCRIPTION:

GATCCCTGGT ATNGTCCGAA TTGTAAAGAA CATCAGCAAG CCACAAAGAA ATTGGATTTA 60
TGGTCCCTGC CTCCAGTACT TGTAGTACAT CTNAAGCNAT TTTNTTACAG TCGATACATG 120
AGAGACAAGT TGGANACCTT AGTTGATTTT CCTATCAATG ACTTGGATAT GTCGGAATTC 180

EP 0 679 716 A1

5 TTAATTAATC CAAATGCAGG TCCTTGCCGC TATAATCTGA TTGCTGTTTC CAACCACTAT 240
 GGAGGGATGG GAGGAGGACA CTATACTGCT TTTGCAAAAN ATAACGATGA TGGAAANTGG 300
 TACTATTTTG ATGACAGTNG TGTCTCCACT GCATCTGANG ACCAAATTGT GTCCAAAGCA 360
 GCATNTGTAC TCTTCTACCA GAGACAAGN 389

10 SEQ ID NO:7345
 SEQUENCE LENGTH:367
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08585
 SEQUENCE DESCRIPTION:
 15 GATCTGTTAC CATCAGGTCA ATTCCTAGTA TGCATAAAATT TTTAACCCT TTAAAAAGAG 60
 ACCTATGTTG AAAACCCCTG AAAATTCCT GAAGAAAAAT CATTACTCTT NTTCTCNGTA 120
 AATCATATCA TCTGAAATAT TACAAATTC AAATTCTAG GTGCTATATT AATTCAATAT 180
 TACAATAACT CTTACCTAAT TATTCTTACA AGTTTAAAGT TGTGGTAGTT TAGTGATTTT 240
 TTTAAAAGAT GTGTGAAATG TTCTCTGCAA AATAATTCAG GCCACTGTCT CCTTTTATAT 300
 20 ATNATNATNN TNGTNTGTGA TGANGACCAG TGAATTACGA TATTTAAAGT GAGAGACCTT 360
 AATTNTN 367

25 SEQ ID NO:7346
 SEQUENCE LENGTH:23
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08586
 SEQUENCE DESCRIPTION:
 30 GATCCCGTCT ATAATCNAAC AAA 23

35 SEQ ID NO:7347
 SEQUENCE LENGTH:103
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08587
 SEQUENCE DESCRIPTION:
 40 GATCGGCTCC CTGTCGCCGCC CGAGGAGGGC TGGACCTTTC GTGTCGNACC CTTGGGNGCG 60
 NNGAGACTGG GTGGGGAGGG TGTGAATAA AAGGGAAAAT AAA 103

45 SEQ ID NO:7348
 SEQUENCE LENGTH:367
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08588
 SEQUENCE DESCRIPTION:
 50 GATCTGGTGG CTGAAAAAC AAGAGTGACA TTAACAAATC ACAAAAAATC AGTTAGGGCT 60
 GTGTTTTTAC ATCCAAGACA TTACACATTT GCATCTGGTT CTCCAGATAA CATAAAGCAG 120
 TGGANATTCC CTGATGGANG TTTATACAA AATCTTTCCG GTCATAATGC TATTATTAAC 180

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EP 0 679 716 A1

5 ACATTGACGG TAAATNCTGA TGGAGTGCTT GTATCTGGAG ACAGAAGAAA CTCATCCAGT 240
CAGCTGGAAA CCAGAAATTA TCAAGNGAAA GNGATTTTAA TGAATGTGGA ATTTTTCCTC 300
TCTCTTTTTT TTTCTTTTAA ATTAAANAA AAAAGGCTTG GNGTTCATGG GGGNTTCCC 360
GNCCTTN 367

10 SEQ ID NO:7349
SEQUENCE LENGTH:22
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08589
SEQUENCE DESCRIPTION:
GATCCCGCCT NTTCTTACA AA 22

15 SEQ ID NO:7350
SEQUENCE LENGTH:378
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
20 CLONE:HUMGS08591
SEQUENCE DESCRIPTION:
GATCTTTTAA AAATTCATCC CATATCCAGA AAGTACCAGT TATAAAGATT GCTGACCAAG 60
CAAAGTTTGG CATCAAAGTG TCACCTCATT GCTCTGACCA AAGACTGACT GTTGTGGTTT 120
25 TAACTCCTCT CTGTAAAGCA TTTTGCATTT TCCCAAGCT CCTTTCTGAA AGAAGAACCA 180
GTGCAGAGCG GCCTTTACTT TCAATTTCTA CTGCTGAATA GACTACTTAG AGAAAATGTG 240
AGTTTCAGTG TGAACAGAAAT GGATTAGGAT GACGAGTTTG ATGGGCATTT TCAGTACTGT 300
ATCTAAGAAA AAAAAATTNG CACAGCTAGG AGCCTCTGAC CATTGTCTGG GTGTTTACG 360
30 TGGGTCTGT TCATCAA 378

35 SEQ ID NO:7351
SEQUENCE LENGTH:379
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08592
SEQUENCE DESCRIPTION:
GATCAAAGTC AAAAATAGCA ATGNCTCCCT ATCCCTCACA CATCCAGACA TCATGAATTT 60
40 TACATGGTAC TCTTGTGAG TTCTATAGAG CCTTCTGATG TCTCTAAAGC ACTACCGNTT 120
CTTTGGAGTT GTCACATCAG ATAAGACATA TCTNTAATTC CANCCATAAN TCCAGTTCTA 180
CTATGGCTGA GTTCTGGTCA AAGAAAGAA GTTTAGAAGC NNNNNCACAA AGGGTTGGGA 240
GCTGATGAAA CTCACAAATG ATGGTAGGAA GAAGCTCTCG ACAATACCCG TTGGCAAGGA 300
GTCTGCCTCC ATGCTGCAGT GTTCGAGTGG ATTGTAGGTG CAAGATGGGA AAGGNTTGTA 360
45 GGGTGCAAGC TGTCCCAAN 379

50 SEQ ID NO:7352
SEQUENCE LENGTH:119
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08593

SEQUENCE DESCRIPTION:

GATCTGCTTT AGTGAGAGGA CAATTTCTGA TTGATTGTTT TCTCTTCAGG CCATCTCACC 60
TCTTCATTCT CTTGTNACAT TTGAAGCAGT TGATATAATG GGTTTATANN NNTAAAAGN 119

SEQ ID NO:7353

SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08594

SEQUENCE DESCRIPTION:

GATCAGCTAT TTTCAACATA ACTGAAGGCA TATGCTGGCC CATAAACACC CTGTAGGTTT 60
TTGATATTTA TAATAAAATT GGTGTTTTGT AAA 93

SEQ ID NO:7354

SEQUENCE LENGTH:297

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08595

SEQUENCE DESCRIPTION:

GATCTCTGTA GACTTGCCTC TTCTGGACAT GACATAGAGA AAGGAGTCAT AAATTCTCCA 60
AGGTGTCTGT TTCTTCTTTA ATGTCATTCC CTGTTTCTCC TCACATTCCC TCCCCATTTC 120
CTGGGCCCCAG TCTCACACTG GTCCTTGCTT ACCCTAAATG CTATTAATTC CATCACTCTG 180
AGTATGGTGT TTGCTGTCCG CTGAATGCCA AGAGCTTCAA GAGTGTGTGT AAATAAAGCC 240
ACACCTTTAT TTTTGTATTA TTCTGAACCA TGGCTAATAA ATTGTTTCAC CAAGAAA 297

SEQ ID NO:7355

SEQUENCE LENGTH:45

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08596

SEQUENCE DESCRIPTION:

GATCTCCGAT GTGATGAATA CGAATAAAAG GCCCTTAATG GCAAA 45

SEQ ID NO:7356

SEQUENCE LENGTH:212

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08597

SEQUENCE DESCRIPTION:

GATCTGTTCA TGCTTTTGTG TTCGTCACTG CGGCGGGGCC CTTTGATGTC TTCATCTGTA 60
TGGGGTGGAA AAATCACCAG GAATCCCCCT NNNAGTTCTT TGAAAAAGTT CCATGACTCG 120
AATATCTGAA ATNAAGAAAA CAAACCGACT CACAAACCTC CAAGTAGCTC CAAATGCAAT 180
TTTTAAATG GAAAACAAAA ATCTGAAAGA AA 212

EP 0 679 716 A1

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SEQ ID NO:7357
SEQUENCE LENGTH:250
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08598
SEQUENCE DESCRIPTION:
GATCTAAACT GTATTTTCCA ATTTAAATTA AAAATGTAAT ATAGATTCAG AAAGGTTTCAT 60
ATTTTCTAA TGACTTCATT CTATATTATT TTGTTAGGTT GCATAAAGAA GCAAGGAATT 120
GTACTTGTAT TAAAAGATGA GGAAAGCTAT TAGGNTTATT GGNCCATGCC TGAAATNGGG 180
CCTTTGCCCG TAAAAGGAA AGGTTGGCCC NTTTTTNAAG GGGGCNTTNA TNTGNNTTNN 240
TNTTAACCAN 250

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SEQ ID NO:7358
SEQUENCE LENGTH:112
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08599
SEQUENCE DESCRIPTION:
GATCTGGCTT NGCCCTAAGC NNCTTAGGNA CAGAAGAGTC CATGGGGGCA TGGCACAGAG 60
CTGGTCCTGT ATTCTCCAGG GTCCGGAGCT GGCCAGGGGC GGGGAGGAGG NN 112

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SEQ ID NO:7359
SEQUENCE LENGTH:386
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08600
SEQUENCE DESCRIPTION:
GATCTTATTT NACTTCATCC TTCTTGTAAC CATATTGTNA ATCAGGAAAG CTCAATCAGA 60
TACAAAAACA GGAAATTCGT TAAAAGTAAC GTTTGTAGAA GATACTAGTT GAGGAGGAGA 120
GACTTTTNG AGAGCCTGTG TGTGTTTGT AACTCTTGT GACCGTGCCA AGAAAACGTA 180
AGCAAAATATC AAGCCTCCAA ATCCTAATGC AACANGTCCT GANTCCCTTG CAGGTATGTC 240
CTGGATGATG ANTACACAAG CTCAGTAGGC TCCAAATTC CAGTCCGGTG GTCCCCACCG 300
GANGTCCTGA TGTATAGCAA GTTCAGCAGC AAATNTGACA TTTGGGCTTT TGGGGTTTTG 360
ATGTGGGAAA TTTACTNCCT GGGGAN 386

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SEQ ID NO:7360
SEQUENCE LENGTH:341
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08601
SEQUENCE DESCRIPTION:
GATCCGGGNA TTATACAGGT ACTGTTGGAA GTATCTTGGG GATTTTCCTG ATAAGAACAG 60
TAGTGATTGN ATAAAAGGA CAGGATGTAA AGTGAAATCA GTAAAATATC TTAGTAGACA 120
GAGGGTGCTG AAATTTTAA ACATGTGTAA AAAGTTCTTC CTATGCATTA ATTTTCAGA 180
TACCCTTAAA ATGTTTAAGG AATGTAATTC AAAATACTGT TTAANAGAGA CATGTGACCA 240

TCATTCTCCC AGCGAATGTG AATCATTTAG TGNCTACTC AAAATTAGGN GTAAATGTAT 300
 ATGTACACTA TAAGANTAAN ANTCGNTACC ATTTCTTTAA A 341

5 SEQ ID NO:7361
 SEQUENCE LENGTH:92
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08602
 10 SEQUENCE DESCRIPTION:
 GATCCGTCAG NCGACTCTAA AGCTAATTGT TTGTATTGGC ATGGNTTTTG CCAGGCATCC 60
 AAAATTCAGA TTAACAACC AATGCAATCA AA 92

15 SEQ ID NO:7362
 SEQUENCE LENGTH:359
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08603
 20 SEQUENCE DESCRIPTION:
 GATCTCGTCT GAAACAGTGT TTGGAAGTGG GAACAGTTTT GTCCTGTATG CTGATGTGTC 60
 CAGAATTTCA TTTAATGATA GACGGAAAAT GTGTGGTTAC TGAAACTGT ATATGATACA 120
 GAATTTTCATA AGAGCCATGC TGTGGGCAA AGCAACTCTT TTTCAACCAC TGCTCATCAG 180
 25 TTTCTGTAGA GACAAAACT CTGTACATAT TTTGGAATCT GAAGAATCCT ATGTAAATCA 240
 TTTGTTACTT AAGTCTGTGA AAAACATATT TCTTTGGAGG AAAATGTATG CATTATAAG 300
 TGTTCATGG AATCAGTTTT TATTGTATCG ATATAATTGT CTCTAAGTGT TGACTGTCN 359

30 SEQ ID NO:7363
 SEQUENCE LENGTH:273
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08604
 35 SEQUENCE DESCRIPTION:
 GATCTCCTTA CACTGCTATT CCAGAGGAGA AGNCACCAGG TCACTCAACT CTTAGTATCA 60
 TCTACTGGAA ACTGGCTAAG ACAGTATTTA TGTGCTTCTC TCACAATAAC AGGAAGATAN 120
 CTATATCACA CAATGGATAT GGCAGTTGGC TTGACTGCAG TAATCATTTA ACTATGTATA 180
 TGTATATCAA AACATCATGT TGCATACCTT AAATATGTAC AATAAAAAATA AAATCTAAAG 240
 40 GANATAANTT AANATTACCT TCCACCAGGC AAA 273

45 SEQ ID NO:7364
 SEQUENCE LENGTH:358
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08605
 SEQUENCE DESCRIPTION:
 GATCAGGGAT GCAGTACCAC TTGCAGTCAA ATGAATTCCT TCGAAATGTA TTTNNACTTG 60
 50 GACCCCCAGT GATGCTTGAT GCTGCAACGC TAAAACGAT GAAGATTCTT CGTTTCGAAA 120

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GGCATTATA TAACCTGCA GCCTTCAAAG CTCGAACCAA AGCTAGAAGC AAATGTCGAG 180
 ATAAGAGAGC AGATGTTGGA GAATTCTTCT AGATTTTCAG AACTTGAAGA CTATTTTCTA 240
 ATTTCTATTT TTTTCTAN TTCAATGTAT TAAACTCTA GACACAGTTT TTATCCTGGA 300
 TTAACCTAGA TAACCTTTGT AGCAGTGGTT ATATTGCTTA TAATTTAATG TACAATTN 358

SEQ ID NO:7365
 SEQUENCE LENGTH:263
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08606
 SEQUENCE DESCRIPTION:
 GATCTACNGT GAATATAAAA TGTTATTAGC CTTATCAAAT AAATAAATGA GCANAGCTCT 60
 TTGTTGCGTC TATTGAACTG GGTGATAATN ATTTAGATGT GAATGTTAGG TCAGTTCTAC 120
 TGTAATAACT AATCGTAAAA GTTGTAATN TGTTAATGG ATGAAATGTA CANTTTTTGT 180
 ATTCNTACA GCTTTGTCTT TTTTAAATAT GAAGTACTTA AATGTACTTA CTATGATAGG 240
 AAATATAATG TGTGCCTTCT AAA 263

SEQ ID NO:7366
 SEQUENCE LENGTH:366
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08607
 SEQUENCE DESCRIPTION:
 GATCTAAGTA AAATGTTTGG AAGTTTATCA GAAACTAAAT GTACTTTTAA AACGTATAGG 60
 GTCAGGGTTG GGGGAAAAAT ACAGGTATAG TAAGNAAGAA AAGTGACCCA TGAAGAAAGC 120
 ATCGTGAGGT TGTATGTTGG TTGACTGTGA TTAAATGCG GGGCTGGTGT AAGTTGTAAG 180
 TGGTGGCTGA TTGCCGTGTA ACTATGTACA TGATTGTTGG GATGGCTGTC CCATATTTTG 240
 TATATTGGAA TAAAAATTC TATAANTTAT NGTAACTAAA AGTAAATATT CTAAATTAAG 300
 TCCCACTNCT TAAGTCACAT GGCTTCTGTC TTGGGAATTT TACCTTTTAA NANGATTATT 360
 TTNAGN 366

SEQ ID NO:7367
 SEQUENCE LENGTH:378
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08608
 SEQUENCE DESCRIPTION:
 GATCTTGGCT GTTCTTTCTA TTCTGCCAGC CATCTTCCAA AAAAACTAA AGCAGAAATT 60
 TGAGTAAAAA TAATCATCTG ATTTAAAGTT TTGCTGTCAT CACCATCTCA GGATTAACAG 120
 CTGCGACTTT AGGTGGGGTA TATTTTCTTC TCCTAAGAGA ATAGACAGTT TTTCCAGATT 180
 CATCATCATT GACTGTCAAG AAAGGACCCT TCAGCAAGGC TGTACCNTCA ATGCAGTTGA 240
 TGGCCTGTNT TCACGGNTTT ACAGACTTGG CCTGATGCCC ATGTAANTTC AAGCTTTGGC 300
 TTGTGGNAAC AACCACANGG ANGCAAGGC ATCTGTNGGT GCGGGNGGGC AAAGCAGGCT 360
 TANTNGGGN GGTGACN 378

EP 0 679 716 A1

SEQ ID NO:7368

SEQUENCE LENGTH:421

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08609

SEQUENCE DESCRIPTION:

GATCTAGAAA CCCTTTTACC TGGCCCCAAG GCTCTAGCCA GTGGTCTCTG GTGACCCCTG 60
 CAAGCTCAGA GATGGGGNTG TTGTGTGCCA GGCATGAGCT TCCCCAAGAG GCAGAGGGCA 120
 TACATCCCAT TTATACATTC CCTAAGCCCA CCCCTGTNTC CTCAGGCACC ACCTGCCCTG 180
 GTCAGCTCCA GCCAGTCCCT GCTGGAGTGG TGCCAGGAAG TNACCACTGG CTACCGTGGG 240
 GTCCGAATCA CCAACTTNAC CACATNNTGG CGCAACGGCT TGGCCTTCTG TGCCATCCTG 300
 CACNGATTNT ACCCAGACAA GATGTGAGCT GCCAGAGGGG TGGGACGAAT GGGGGAGCCA 360
 TCAGGGAGGG CAGTNTGGAC CCAGCTTTGA CCAGANAATT TTTNGAGTGC TTTGCTNGTT 420
 N 421

SEQ ID NO:7369

SEQUENCE LENGTH:189

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08610

SEQUENCE DESCRIPTION:

GATCAGCAAT GAGTCGGCAA TTGACTTCTA CAGGAAGTTT GGCTTTGAGA TTATTGAGAC 60
 AAAGAAGAAC TACTATAAGA GGATAGAGCC CGCAGATGCT CATGTGCTGC AGAAAAACCT 120
 CAAAGTTTCT NCTGGTAAGA CNNCAGNTGT NAAAANGCCA GCAACCTGNC CAATTTCCAN 180
 TGANCCTTN 189

SEQ ID NO:7370

SEQUENCE LENGTH:147

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08611

SEQUENCE DESCRIPTION:

GATCTGGTGG AGAAAAGAAG CAAAACCTTG TTTTAATTGA CTGTTTTTTT AAAATCTTTT 60
 TGGATATTGT TTTGCATCCC ATACATTTGC AACTGGCAAT GAAATGTAAG CTTTGGTTCT 120
 GAAAAAAAT AAATGTAGCT ATTCAAA 147

SEQ ID NO:7371

SEQUENCE LENGTH:108

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08612

SEQUENCE DESCRIPTION:

GATCCACATG CNTGTGGCAG CCCAAGGCTT TGTTGTAGGA GCAATGACTN TTGGTATGGN 60
 CTATNCCATG TATCGGGAAT NCTGGGCAAA ACCTAAGCCT TAGAAGAN 108

EP 0 679 716 A1

SEQ ID NO:7372
 SEQUENCE LENGTH:273
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08613
 SEQUENCE DESCRIPTION:
 GATCTTTGAG CTGTGGTTTC TCCACCTGTG GAATATAAAT GTNAACAACA TAACCTTCTT 60
 TCTAAGGATA ACTTGAGAAT TAACAGTGCA AGTAAATGGA AAGTATAGTA CCTGTNACCA 120
 GATATTTAAT AGTAGTTATT ACGAAATTGT AACTCACTAC CTGTNTTAGT TGAGGTTTAT 180
 CTTCAGCCAT GCAAATCAAA ATCAGATTAT TTNCACATCA GCAGATAAGG TTTGNTGAG 240
 ANTCTNATT TNNTTTGNT ACTCTNATT NNN 273

SEQ ID NO:7373
 SEQUENCE LENGTH:419
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08614
 SEQUENCE DESCRIPTION:
 GATCACAGAT GTAGTTCCTG AGGTTGAGTC TCCTTCTCAG ATGGATGTTG AATTGGTGAG 60
 TGGGTCTCCT GTGGCACTCT CACCCAGCC TCGATGTGTG AGGTCTGGTT GTNAGAAGCC 120
 TCCATTGTG AGTAAGGACT GGGACAATGA ATACTGCAGC AATGAGTGTG TGGTGAAGCA 180
 CTGCAGGGAT GTATTCTTGG CCTGGGTAGC CTCTAGAAAT TCAAACACAG TGGTGTGTTGT 240
 NAAATAGTCC TTCCTGTCTT CCAAGCCAGT GAAGAGTTAT CTGCTGGGAA AGTGTCCAAG 300
 AGCCTGTTTT TGAACACAA GCTGGGCTTC TGGTAGTGCC TCATCACAAC CCATGATGGC 360
 TGTTCATGGT TTCACCCCTT TTNTTCCNT CAGCAGAGGC CAGGCTATTG GAGCAGTTN 419

SEQ ID NO:7374
 SEQUENCE LENGTH:209
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08615
 SEQUENCE DESCRIPTION:
 GATCCTAATT TGCACACTAT CACCTAAAGT CCTAGCATAC CTGCATTTC AGGAAATCTA 60
 ATGAGAAAAG AGTCCTGAAA GCCTCTAATT CCTGGAAACA TATGAAACGG CTAGAAAAAA 120
 GTAGACTATT TTGGAGGATT AAAGACAATG GAAACTTTTT ACATTGTTTG TCTCACATAA 180
 TGAATAAAAT TCCACTTTAA AAAATGAAA 209

SEQ ID NO:7375
 SEQUENCE LENGTH:422
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08616
 SEQUENCE DESCRIPTION:
 GATCCGCAGC TCGAAAAAGA ACAAGCCACA GAANCGGGCT CGCTCGTGCC AGGACACAGC 60
 AGTGTCTTTA AAAAAATCAA AACCAGAAGT TTTATCAGCA GCAGGAAGNN TGTGGGACTC 120

EP 0 679 716 A1

5 TGTCCAAGTT CACCGTCACC ATCAAGCCAC TGGCTGTGGA AGGAGTTTNT CCAACAGGGT 180
 CAGTGTGACA GCCACAATT CAGAAAGCAG CCATCCCGCG TGTGTCCTCAA ACAGCAGTGC 240
 CTGCNTCCCG CTCCACGGAG TGTTCENGAA CACCTCCCTC TGAACATAAT CCGGAGCACC 300
 CCACCCCCAG CGAGTCCAC CTGGACCTNT TCAAAGTCAA AACTCTTCTT TNGGGCCAAC 360
 CAATTNTTCG GGCTTTNGAG GCCTTGAAC TGGGGCACCA AGTTTNTTC CATGGGAANT 420
 TN 422

10 SEQ ID NO:7376
 SEQUENCE LENGTH:413
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08617

15 SEQUENCE DESCRIPTION:
 GATCCCACCA GTTCTGGAC GTGGAGATC TGGGGCATCT CCTTCTTATG CCAAGGGGCG 60
 CTTNNNGTTT TCATGGCTGC CCCTCCAGAC TGGGAGAAAC AAGTAAAAAC CCATTGGGGC 120
 CTCTTGATGT CTGGNATGGC ACGTGGCCCG ACCTCCACAA GCTCCCTCAT GCTTCCTGTC 180
 CCCCCTTAC ACGACAACGG GCCAGACCAC GGGGAAGGACG GTGTTGTNT CTGAGGGAGC 240
 20 TGCTGGCCAC AGTGAACACC CACGTTTATT CCTGCCTGCT CCGGCCAGGA CTGAACCCCT 300
 TCTCCACACC TGAACAGTTG GCTCAAGGGC CACCAGAAGC ATTTNTTAT TATTATTATT 360
 TTTAACCTG GACATGCATT AAAGGGTCTA TTAGCTTCA GAAAAAAAAA ATN 413

25 SEQ ID NO:7377
 SEQUENCE LENGTH:157
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08618

30 SEQUENCE DESCRIPTION:
 GATCCACAAA TGAAGGGAT ATAAAAATAA TGTCTAGGT AAGAAACACA GCAACAATGA 60
 CTTAACCATA TAAATGTGGA GGCTATCAAC AAAGAATGGG CTTGAAACAT TATAAANTT 120
 GACAATGATT TATTAATAT GTTNTCTCAA TTGTAAA 157

35 SEQ ID NO:7378
 SEQUENCE LENGTH:399
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08619

40 SEQUENCE DESCRIPTION:
 GATCCTAACA GCTTGCTTA TTTGGTGCAT CTGCTCAATC AAGTCTAATA GACACAAGGA 60
 TGGCTTTCAT CGGCTCAGGC AGCATCATGA TGAGTATGAA GATGNAATTC GCATGATGTC 120
 45 TACCGGTCC AAGAAGTCCC TCCTAAGCCA TGAGTTCAG GATGAAACAG AACTGAAGA 180
 GGAAACATTA TATTCTAGCA AACATTGGNG GNCACATTT GCATATCTCC CAGCATAAGT 240
 ACCAAGCAAA ATTACAGTTC CTCTGGGAG AACACTGCAT TAAGGATGCG AGACTCTCTT 300
 GCTTCTTCAA AGNGCTTTTG GGAATTTAAA TTGCTAANTA TGTATTCTCA AAANAANNNN 360
 50 NGGGANNTTT GGGGGGNTG GGTTTNCGG AGNGANGNN 399

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SEQ ID NO:7379
 SEQUENCE LENGTH:34
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08620
 SEQUENCE DESCRIPTION:
 GATCGGTCCC TAAACCTCAT TAAATATTT GAAA 34

SEQ ID NO:7380
 SEQUENCE LENGTH:405
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08621
 SEQUENCE DESCRIPTION:
 GATCTNATGG ATATGGCAGT GGACGTGGAT TTGGGGATGG CTATAATGGG NATGGAGGAG 60
 GACCTGGAGG TGGCAATTTT GGAGGTAGCC CCGNTTATGG AGGAGGAAGA GGAGGATATG 120
 GTGGTGGAGG ACCTGGATAT GGCAACCAGG GTGGGGNCTA CGGAGGTNGT TATGACAACT 180
 ATGGAGGAGG AAATNATGGA AGTGGAAATT ACAATNATTT TNGAAATTAT ANCCAGCAAC 240
 CTCTAACTA CGGTCCAATG TNGAGTGGAA ACTTTGGTGG TAGCAGGAAC ATGGGGGNAC 300
 CATATGGTGG AGGAAACTAT GGTCCAGNAG GCAGTNGAGG AAGTNGGGGT TATTGTNGGA 360
 NGGNGCCGAT ACTGAGCTTC CTCCTATTT NNCCATGGCA TTTNN 405

SEQ ID NO:7381
 SEQUENCE LENGTH:261
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08622
 SEQUENCE DESCRIPTION:
 GATCCTAGCC AGATTACCAG AAATTNCGTT AGTCATCTAA ATCAGAGACA TCAATTTNAT 60
 TATGGAGAAT TTNTGAATCT CCAGCTAGAT GANGAAACCC AATACCAAAC TCCTGTTGAA 120
 GAATCTTTNC AAGTAAACAT CTGAAGGCTG TAGACATCTC TCCATCTTG TACCTGCAAG 180
 TCCCATCTTT AAGGGGGAAA CTACATGANG TCACCGTTAC AGTAACTTGA TGTGTATATT 240
 AATAAAAGTA ATTCAGTCAA A 261

SEQ ID NO:7382
 SEQUENCE LENGTH:44
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08623
 SEQUENCE DESCRIPTION:
 GATCTATGTA TAGATAAAGA TTAACATAT CTAGTACAAG GAAA 44

SEQ ID NO:7383
 SEQUENCE LENGTH:109
 SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08624

SEQUENCE DESCRIPTION:

5 GATCTGTTTC TTAAAGCTAC AGGGTTTAAA AAATAAAAAAT GAGTGAAAAAT ACTTGATGTT 60
TCTTGAAAGA TAAATTTAAT AATAATAAAT AAATACATAA ATACATAAA 109

SEQ ID NO:7384

10 SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08625

SEQUENCE DESCRIPTION:

15 GATCATTGTG GCGTCCACAA TGTATATATA AAACAAGTTG TACACATTAC CATGTATAAA 60
TTTGTGTTTG TCGATTATAT GTCAAGAAAG AAA 93

SEQ ID NO:7385

20 SEQUENCE LENGTH:54

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08626

SEQUENCE DESCRIPTION:

25 GATCTTAATC TTATTCTCTG CCATCTTCAA ATTAAAAAAA AATTCTTTG TAAA 54

SEQ ID NO:7386

30 SEQUENCE LENGTH:140

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08627

SEQUENCE DESCRIPTION:

35 GATCCCTTCA GAAGAAGACA TGATAGGATT GTAAGTTTTT NTCTAACTTT GGATGCAGTG 60
AGATGACCAG TGTGTTCCAG TTAAAGAAGA AGAGTGTTTT AAAATCATAA ACCAAATAAA 120
GAATCCTACC TTACATTAAA 140

SEQ ID NO:7387

40 SEQUENCE LENGTH:206

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08628

SEQUENCE DESCRIPTION:

45 GATCCACCCA CCTCAGCCTC CCAAAGTGCT GGGATTACAA GCGTGANCAC TGTGCCTGGC 60
CCTTTTTTTT TTNAAAGAGA TGGCATCTTG CTATGTCGTC CAGGCTGGTN TTGAACTCCT 120
GAGTTCAAGC AGTCCTCCTG NTTCAACATA CAGNTACAGG TACCCCCAC TATACATTTT 180
NAATANGGNT TCATGGNTCA GNGGGN 206

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SEQ ID NO:7388

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EP 0 679 716 A1

SEQUENCE LENGTH:130

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08629

SEQUENCE DESCRIPTION:

GATCCCATTTG CTTTCCTGTT TTAATAATAT TTTATGCTCT TTATTTCCAC TTCTGTGAAT 60
GTGATATTTC TATTTTNTGA TTATGTTACT GAATAAACAA ACTTGCTACA TAAATNCTT 120
AGCAATTAAA 130

SEQ ID NO:7389

SEQUENCE LENGTH:355

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08630

SEQUENCE DESCRIPTION:

GATCCAATTG TGGATTTATT TACAAACATC AAATGCCTTC AAGCCAATCC TTTTTGCTGT 60
ATGTTTTGCA GCCTACTGTA GTAGATACGC AACAGATAAT GTGGGAAAAA AAGAGATAAG 120
AGGAGGAAGC TAATAAGAGA CTGTCAAGAT TGTATACCTT CTGGGTTTCT TTAAAGAATT 180
TGTTGCCTTT NTACTATNNC AGCAAAGCAG CATTGTTGTTA CTGACTGCCT AAAATCACTT 240
AATCTCAGGT GAACGCATCA CTTGCCAAAC TGTGGGAATG CTATTTGTGT TTTGTTGCAC 300
TGTTTTTTTC GTTTGTTTGT TTGTTTATTT GGTGGGCTTT TTGGAGAGGG AAATN 355

SEQ ID NO:7390

SEQUENCE LENGTH:96

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08631

SEQUENCE DESCRIPTION:

GATCTAAAAT NAGAGAAGCA TTTAAAAAAT NATTTCTGTC TTGGTTCATC TTATTGTTAG 60
TGGTTTATGC TAATAAAAAT TGATTTAACA ATCAAA 96

SEQ ID NO:7391

SEQUENCE LENGTH:411

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08632

SEQUENCE DESCRIPTION:

GATCCAGATA TGTCTGTTAT GAAAGATATC AGTATAGGTA AAGCCACAGG CAGAGGTCAG 60
TACTGATAAT TAATGTAGTA TAAATACATC ATTTACCATT TTATTTTAAA TAGGAAGCCA 120
TCAAGCATGC TAGAATTGTG AACTTTTCATT ATATNTTTTT GTTGTGACA TGAATTAACC 180
TGGCCAAAAA CAAAAAAGAA NAAAAAACC ATGCCATTTG TCATGTAAAC CTTTTTTTTA 240
TCCCTATGGG CCTTNGGGGN CNGAATCAGT ACTTCAGTTA TTGTAATNGT GNGCTAACCT 300
CAAAATTTN TNACCCNGTT GCCCTTTTCA TGNCTAACCT AATTNTCTGG GGGNTTGGTA 360
TGTTTCNCCC GGTCTGNTC ATGTTTACCG NNCTCTTTT TNGNGGGTCC N 411

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SEQ ID NO:7392
SEQUENCE LENGTH:248
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08633
SEQUENCE DESCRIPTION:
GATCTAAGTT CTAAGCCAAA TAACTAACAA ATTCCAGGTA CTATGCCATC ATATAATATT 60
TATAAATGTG TTATATATAT TATGAGATAT GTATAAGAAT ATAGAATGAA TATGTAATAA 120
ATTATACTTT TAAATTAATA TTTTCAAATA TATAAATATG TAGTTTCAGA CTGAAACTGT 180
TCTAAACTAT ATAAATGTAT TATGTATGTT TTATAAATA CAGTACTCAT AATATGTATT 240
ATATGAAA 248

SEQ ID NO:7393
SEQUENCE LENGTH:247
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08634
SEQUENCE DESCRIPTION:
GATCCTTCCA CCTCAGCCTC CCCAGTAGCT GGGACTACGG GGTGCCCCACT GCACCCAGCC 60
TGTGTCTTCC AGCTCTAATC AGAAAGCTGC TTGTCGAGGG CATGGGGAAT CAAACTGAAT 120
GAACTTTTCT CTGCACTGTG GCAAACTGT TATTTTATG GATTTTACTA AATGGCGTTA 180
CCTTTTCAAG ATTTATATGT TTGTATAATC ATAAGAAAAT TGAGCCATTA AAGCCTTGTT 240
ATTCAAA 247

SEQ ID NO:7394
SEQUENCE LENGTH:108
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08635
SEQUENCE DESCRIPTION:
GATCCAGGCT CGTAGNTCAC TCCCTGCCCC TNTCCCAGAG ATGCTTCACC ANCACCTGCC 60
TCTGAGACCT CGCTCTCTGT TCCAGCAACC CTGGNTTGGG GGGTCANN 108

SEQ ID NO:7395
SEQUENCE LENGTH:61
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08636
SEQUENCE DESCRIPTION:
GATCTTGAGC TCCTGGCCTC CAGAATTGCA GGAGAATAAA TNTGTGTTGT TTTAATGAA 60
A 61

SEQ ID NO:7396
SEQUENCE LENGTH:275
SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08637

SEQUENCE DESCRIPTION:

5 GATCAGTGTT TGNTTGGGAC TGAAGGGATG ACTAATAAGG AGCAGAGGAA AGCTTTTGGG 60
 GATGATGGCA CTCTTCTCTG TCTCCATTAC GGCATTGGAT ACAAGTCTCT AGAACTCCCA 120
 GAACTCGTCA CTGAAAGGGG TGAATAGTAC TGTATGTAAA TTAGGTCCNN ATTTTNAAT 180
 GGGANAAATA AATATAAGA CAAATTACAC TGCTGTATTA NNTCANNTGA TAGATGTAAA 240
 10 AGTCTAATAT ATACGTNTTT TTCNTGATAN NCCCN 275

SEQ ID NO:7397

SEQUENCE LENGTH:288

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08638

SEQUENCE DESCRIPTION:

20 GATCCACGCC CTTCCCGCCA GTCTGGAGAG GGCCCTGACC GAGGACTCAA CCCAAACCAG 60
 TGACACAGCT ACCAATTCTA CTTTACCTTC TGCAGAGGTG GCGTTACAGG CAAAGTGAGG 120
 AGGGAGCTGG GGGACACTTT CGAGCTCCCA GCTCCAGCTT CGTCTCACCT TGAGTTAGGC 180
 TGAGCCACAG GCATTTCCTG CTTATTTTAG GATTACCCAC TCATCAGAAA AAAAAAAG 240
 GCCTTTGTGN CCCNTGTTT GGGGGGATTA ACCNGTTTGG GGTTTAAA 288

25 SEQ ID NO:7398

SEQUENCE LENGTH:290

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

30 CLONE:HUMGS08639

SEQUENCE DESCRIPTION:

35 GATCTAATTN CTATTTCAAT TATTCAGAAA TAGCATTTAG ACATAAAANC CAATGTCTCA 60
 CTTTGTAATA TAACCTTTGG CTAATTTACA CACATCTAAT ACAGCGTGTT ATATAAGTTT 120
 TAAGTAATAC AATGAGTCAC TACTATCATT CAGTTTAAAN TATTTNNGT GTTAACAGGG 180
 CTGAGAATAT CATGTGGTTC AGTCTTCTGN NGGNCGTTAT ATNATACCNG CNTAGTGCCT 240
 TTGACCATGC NGAATATCCT CAAGGCCAGA CATCCTACAN CGGAACTGNN 290

40 SEQ ID NO:7399

SEQUENCE LENGTH:267

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08640

SEQUENCE DESCRIPTION:

45 GATCCNTAAG TACTCTTCTC TANAGAATAA AATGCATTAC CTAGGCTTTA ATAGTTAAAA 60
 GCTGGNATTT ATNGACCCGC ATCTACTTGA TTTCANAATA TGCATTGATT TTAGTATGTT 120
 GCTGNTCTGT TATCTGTACT CAACTATANC AGAGTAATAT ATTGTGGTNT TCTGGACCAA 180
 AANTCGTGCT ACACATATGT CTGATGGCCT TNACCTTTTC ANGACAACAG TTGCAATTTT 240
 50 GAGCACAGTC CNCCTAGTGG TANTCTN 267

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EP 0 679 716 A1

SEQ ID NO:7400
 SEQUENCE LENGTH:127
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08641
 SEQUENCE DESCRIPTION:
 GATCGGGAGC AGCCCCACTG GACCTAGGTG CCCCATCTGT TGGTCATCCA TCCTGAAGGG 60
 10 ACAGGAAACC TCCCAGGCAG TTATTTTTTT TTCTCTATAT TTCTAGTAAA GTTTTCGATA 120
 TGTAAAA 127

SEQ ID NO:7401
 SEQUENCE LENGTH:168
 15 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08642
 SEQUENCE DESCRIPTION:
 GATCACTAGT GCAGGAGAAC ATTACATTTT CTTCTGAAGG CAAAATGCTT GTAGGTTTTG 60
 20 CCTCTACTTT GTATTACTT TAAAAATTGC ACTTGTTTAC CTACCAGTGT TTACGAAATC 120
 CTGTATTGG GATGCTTTTT CTATAATAA ATATTATAAT TTGTGAAA 168

SEQ ID NO:7402
 25 SEQUENCE LENGTH:323
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08643
 SEQUENCE DESCRIPTION:
 30 GATCATAGTA AAGAGTAGTC AATAGGGTCT TCAGCTATTA ATTGTAGAGG TGATTAAAC 60
 CAACAAGGAG TTTATGTGTC AAAGGAGATA AGGAATGAAT ATAAAGATTG CTATTGGGGT 120
 GGCTCTTATT AAAGTGTGTA TTTGTACTT ATCACTACAC GTATCCCCCA AATGCTTACA 180
 TGGGAGTTTG AGGTAGTAT TTTCACTTCC TTGGTGTTAG TACTCTATTC ACATTCTTAT 240
 35 TGTAACCTTC CTCATTTTAC AGATAAGGAA TCTTTGGGGA TTAACCAACC TCCTTTCTGT 300
 AATGGTAATC ATTAAANTAA GTN 323

SEQ ID NO:7403
 40 SEQUENCE LENGTH:94
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08644
 SEQUENCE DESCRIPTION:
 45 GATCACTTAN GCCCAGGAGG CAGAGGTCAC AATGAGCCGA AATTGTGCCA ACTGCACTCC 60
 AGCCTGGGCA ACAGAGGAAG ACTCTTCACA GAAA 94

SEQ ID NO:7404
 50 SEQUENCE LENGTH:226
 SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

5 TOPOLOGY:linear
 CLONE:HUMGS08645
 SEQUENCE DESCRIPTION:
 GATCGGGGCT TTGTGTTTAA CCTCATCAGA CATTATTGCA GCCAGCTGTC AGCCAAGCTC 60
 AGTAACCTTC CAACGCTCAT TTCCATGAGG CTAGAGITCC TGAGAATCCT CTGTAGCCAT 120
 GAGCATTACC TCAATCTGAA CCTTTTTTTT ATGAATGCTG ATACTGNTCC NACATCTCCT 180
 TGTCTTCCA TATCTTCCCA GGTAAATAAAA GAATTATTTA ACTAAA 226

10 SEQ ID NO:7405
 SEQUENCE LENGTH:124
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

15 CLONE:HUMGS08646
 SEQUENCE DESCRIPTION:
 GATCCAGTCA GAGCCAGGGA GAGTGTGTC CTTAAGGAGG GACTCCCTTG TCCAGGATGG 60
 GGCTGGCACT AGGGATGGGG TGAATCTTAG GACAGGAGTG GGGGTCAGAA ATGGGAGGGG 120
 TAAA 124

20 SEQ ID NO:7406
 SEQUENCE LENGTH:206
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

25 CLONE:HUMGS08647
 SEQUENCE DESCRIPTION:
 GATCGACCAG CAGGTGGTCT ATATCCATCC TTCCAGTGCC CTCTTCAACA GACAGCCAGA 60
 ATGCCCAAAG AACTTCCTCC CTNTGGTTGC TGTAGGTGTC TCATTGGAGC AGTGCCTCTC 120
 30 CAAGTTTGAG GATNTGAACA AAGAACTGGG ACTGGTGACT TGTNANNGNN NAGTTCAGAG 180
 GGCAGAGGNC CATCATCTCA GCTTNN 206

35 SEQ ID NO:7407
 SEQUENCE LENGTH:103
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

40 CLONE:HUMGS08648
 SEQUENCE DESCRIPTION:
 GATCCATAAG GAGGGCTGTG TGGNGAAGAT TCAGGGACTG ANTAGAGGGA AAATCGTGCT 60
 ATNCGGTAGC TACAGCAGCC CTGGGAATNC CNTTTTGTCT AGN 103

45 SEQ ID NO:7408
 SEQUENCE LENGTH:116
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

50 CLONE:HUMGS08649
 SEQUENCE DESCRIPTION:
 GATCCATAAG NAGGGCTGTG TGGAGAAGAT TCGGGNCTGG CTGAGGAAAA ANCGGCTGGT 60

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GGTAGCTACA GCANCCCTTG GAATACCCCTT TTGTCGAGGT TTTGGGNATT GTCTCN 116

SEQ ID NO:7409

SEQUENCE LENGTH:60

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08650

SEQUENCE DESCRIPTION:

GATCTATCAT GTATCTTGGT GATTACAGTT AATAATATAT TGTATATTG ANAATNGAAA 60

SEQ ID NO:7410

SEQUENCE LENGTH:91

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08651

SEQUENCE DESCRIPTION:

GATCGCACCA CCGCACTCCA GCCTGGGCAA CAGAGCAAGA CTGTTTCAAA ATAAATAAAT 60
AAAATAATAA ATAANNCTTT CCAACTNCAA A 91

SEQ ID NO:7411

SEQUENCE LENGTH:240

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08652

SEQUENCE DESCRIPTION:

GATCTTGTGG CTAAACCAGC ATTTCTGTGT TTGAGAGATT TCCTGTTAGG TGCTTCGTCT 60
GAAAGTGAAC TCTCATAATT CAAATTGTAT AAANTAAAGC TACATTTCTA AGAGCTTGGT 120
GTAGGGCAAT TGGAATANTG TCCTGTTAGA TAAACAGACA TTTAGCANTC CTGACATTAA 180
AAGGAAATGT ATTTCTATAC AGGATTATTA GCTGTANTAC ANGATATTTA TTTANCCAAA 240

SEQ ID NO:7412

SEQUENCE LENGTH:247

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08653

SEQUENCE DESCRIPTION:

GATCGCACCA CTGAACTCCA GCCCGNATGA CAGAGTGAGA CCCTGTTTCT AAAAACAAAA 60
CAAACAAACA AACAAACAAC GTATCTTACT GACTAACTG GAATGGAAAT CTTATAATNA 120
GTGTTGTATT TTTCTGCAA ATAGTGAATT NGNACTTGGG AGACTTCNNN NCTTTGTTGA 180
AAAAAATTGC CTTTGTGTTG AAATTATATC ACCTGTGTTC TGANCTAAAA TGNTGGGAGT 240
TCATNNN 247

SEQ ID NO:7413

SEQUENCE LENGTH:128

SEQUENCE TYPE:nucleic acid

EP 0 679 716 A1

TOPOLOGY:linear

CLONE:HUMGS08654

SEQUENCE DESCRIPTION:

5 GATCCATAAG GAGGGCTGTG TGGAGAAGAT TCAGGNANCT GACTGAGGAA AAATNTGCTG 60
GTGGTAGNTA CAGCANCCCT TGGAATACCC TTTTGTGCGAA AAATAAAGGA ATTGTCTTCA 120
NCCTGCTN 128

SEQ ID NO:7414

SEQUENCE LENGTH:118

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08655

15 SEQUENCE DESCRIPTION:

GATCCATAAG NAGGGCTGTG TGGAGAAGAT TCAGGGAGCT GACTANAGGA AAAATGTGCT 60
GGTGGNANAT ACAGCAACCC TTGAATANC CTTTGTGCGA GAATAAAGGA ATTGTCTN 118

SEQ ID NO:7415

SEQUENCE LENGTH:151

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08656

25 SEQUENCE DESCRIPTION:

GATCTTTNTN ATATATGCAG TCCATCCCT TCTGTGCCAC TCAATGCCAT CCAGACATGG 60
TTTNCCTC CANGGGCCTT TCTCTCCAGA GGGCACTTCG GCTGCCCTCTG CTTCTCTCA 120
TTCGAGGCCC GNCTCTTCT NACAGAATAG N 151

SEQ ID NO:7416

SEQUENCE LENGTH:374

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35 CLONE:HUMGS08657

SEQUENCE DESCRIPTION:

30 GATCCCGTTG CCTCATGGAA CTCATGCATT TCACGCTGCT TGCCCTTTCT GTGCTACACA 60
GCTGGTTGGG GNGCAAACT GCATCAAATT AATTNCCAA GGTCCAATTG ACTGACGCCC 120
40 TTGACAGCCA TCTACGACTT TATTAACAGG TTAGTGTA GATTTNCCA CTAACCTAG 180
ATTTNCCTT TTTGTAATGC TGTTTATCAG AGGCGGGTGA CANGGGCTGG AAATAAAGNG 240
AGGGGACATG GTGATGAAAC ATGGCAGGNG TGTACAGATA CCAGTGGTGT GTTGCAATGCT 300
CAANACAGCA GCGTCGTCAA TGGCGTCTGC TTGNTTGNNC CATNATGTCT TTGTCATAAT 360
TGGNAANNA NNNN 374

SEQ ID NO:7417

SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

50 CLONE:HUMGS08658

55

SEQUENCE DESCRIPTION:

GATCTTTATG TCACTAGTTA TAGTTAAGTT CATTAGACA TAATTATATA AAAACTACGT 60
GGATGTACCG TCATTGAGG ACTTGCTTAC TAAACTACA AAATTCAAA 110

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SEQ ID NO:7418

SEQUENCE LENGTH:232

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

10

CLONE:HUMGS08659

SEQUENCE DESCRIPTION:

GATCTTTTAA ATTCAGTACA ATCTCTGAAG GTGTAAACAC TAATNTTTTA CAAGAATGTA 60
GAGGCGACTC GGTGCTATGA AGCGTGTTAA ACAACTCACA CAATTGCTAC AAAACACCAG 120
GGAGGGGCTT TTTGTGTTTT TAATTTTAA AACATTTTTC CTTTCTTGT GAGCCATAGT 180
AGGCCTAGAG AGATTACTGG GTTTTATAA TTAAACATTT ATTTCGGTGA AA 232

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SEQ ID NO:7419

SEQUENCE LENGTH:370

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

20

CLONE:HUMGS08660

SEQUENCE DESCRIPTION:

GATCTTAACC AGTCGAGTGG AGTGACATT GTCTGAATAC AGGATGCACA ATGTTGTCAA 60
TCCTGGAAAT GGTCTTCTT TTTGTAAGA TATGTGAATG AAGTGTGGT GTCCTCACCA 120
AGAGGTGGCA CCTAAGGGT CTGAGGAAAT AAATGTATAG ACCCTTATGT ACAGACCTGT 180
GTATAANCAN CTNNGTATA TACATATAAG GATAGCTTTT TTGACCTATA CAGCTGTNCA 240
TAAAGTAGC TGATATTAGT TAGGCCTGTG TCACCAGTTT GGNTTTTTTN CACTGGTNCA 300
TTTGGGATTT NNTTTTNGGT GGTTTAAAT GGCATATGCT AAGGTGTGTG ATTGGGAANA 360
AANNANNANN 370

30

SEQ ID NO:7420

SEQUENCE LENGTH:83

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

35

CLONE:HUMGS08661

SEQUENCE DESCRIPTION:

GATCTGTGTT TTAATGAGTT TCACAGTGTG ATTTTGATTA TNAATTGTGC AAGCTTTTCC 60
TAATAAACGT GGAGAATCAC AAA 83

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SEQ ID NO:7421

SEQUENCE LENGTH:394

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

45

CLONE:HUMGS08662

SEQUENCE DESCRIPTION:

GATCCTCTCT TACCCGCTCC CCAGGTTTGA AACACATAGC CTCATTTCAA GGTGTAGCCA 60

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EP 0 679 716 A1

GGTCCCCCG ACTTCTCT GGGATATAAA AAAGGGGGTA AGGGGGCAA GAGAGCCCTC 120
 TGGGACTCTC CTCCCATACA CACTACACTG CCCCTTCTCC CCCCATCAA ACGCTCAGNG 180
 ACGTTGTGAT GATGCGACTG AGGATTATGC AACGTGGTCC AACCGGAGCG GNCAGCATGA 240
 CCAGCTGTCC AGGGGCTGNC TNCTGCCTTT TCTTTGTAA AAGCCANGAC CCTTGGGGAG 300
 TTTTAATNCT GTTTGTACT TGNCTGTGG GGNCTNCACT TGCTTTTCT ATGGGAGACA 360
 CTTTNAAT TTAACAAGNT GNGGGTTTT TGNN 394

SEQ ID NO:7422
 SEQUENCE LENGTH:134
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08663
 SEQUENCE DESCRIPTION:
 GATCCCCATG CCGAGGGCC CCTCGGCTGG CCTGGGCATG TGATGGCTCC TCACTGGGAG 60
 CCTGTGGGGG AGGCTCAGGT GTCTGGAGGG GGTGTGTGCC TGATAACGTA ATAACACCAG 120
 TGGAGACTTG CAAA 134

SEQ ID NO:7423
 SEQUENCE LENGTH:69
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08664
 SEQUENCE DESCRIPTION:
 GATCATGNGT TTCCTCATGA GGACAAATAC CTGATTTTGA ATAAAGCAGC ATTCAGTTGA 60
 AATAACAAA 69

SEQ ID NO:7424
 SEQUENCE LENGTH:52
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08665
 SEQUENCE DESCRIPTION:
 GATCAACATT TTTTAAAAA GAGCATTAAA GAAAGAATTG TGGTACAATA AA 52

SEQ ID NO:7425
 SEQUENCE LENGTH:356
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08666
 SEQUENCE DESCRIPTION:
 GATCACACAT TCAGACTGTT GTGTCTGTGG AGTTTATAGGA GTGGGGGGTG ACCTTTCTGG 60
 TCTTTGCACT TCCATCTCT GCGACTTCCA TCTGGCATCC CACGCGTTGT CCCCTGCACT 120
 TCTGGAAGGC ACAGGGTGCT GCTGCCTCCT GGTCTTTGCC TTTGCTGGGC CTCTGTGCA 180
 GGACGCTCAG CCTCAGGCT CAGAAGGTGC CAGTCCGGTC CCAGGTCCCT TGTCCCTTCC 240
 ACAGAGGCCT TCCTAGAAGA TGCATCTAGA GTGTCAGCCT TATCAGTGTT TAAGATTTTC 300

NTTNTATTTT TAATTTTTTT GAGACAGAAT CTCACTCTCT CGNCCAGGCT GGAGTN 356

SEQ ID NO:7426

SEQUENCE LENGTH:246

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08667

SEQUENCE DESCRIPTION:

GATCGTCGCC CCTCTGCCAA CTGCGACCCC TTTTCGGTGA CAGAAGCCCT CATCCGCACG 60
TGTCTTCTCA ATNAAACCGG CGATGAGCCC TTCNAGTACA AAAATTAAGT GGACTAGACC 120
TCCAGCTGTT GAGCCCCTCC TAGTTCTTCA TCCCACTCCA ACTCTTCCCC CTCTCCCAGT 180
TGTCCCGATT GTAACCTAAA GGGTGGAATA TCAAGGTCGT TTTTTCATT CCATGTGCCC 240
AGTAAA 246

SEQ ID NO:7427

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08668

SEQUENCE DESCRIPTION:

GATCCTTGCA GGAACCTAAG GATTACAGCA GCTGCAGATG GCAGGAGCTA CAAGCCAGCA 60
GCAGCCAATG CTCANTGGGG TACAAATGGC TCAGGCAGGT CAACCAGGGA AAATGCCAAG 120
TGAATAAAAA ACCAACATCA AGTCGGCTTC CATGCATCCC TACCAGCGGT GAGTGTGGCT 180
GGCAACCTCG ACTCCCTGGT GCTCTTTGCA GAGTTGGGCA GTGAAATNNN NNNNTGCTCA 240
AGGCTCACCT AGGATGGGTA CAATAAAAAG AACATGGGCT TTCAAA 286

SEQ ID NO:7428

SEQUENCE LENGTH:177

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08669

SEQUENCE DESCRIPTION:

GATCACTAAC AGGTGATGGG CTTTGTGCCC ACTCCAGAGA TATTGTGGGA GACAAATTCT 60
TTTAACAGCC TGTCTCCCG GCATCAGGAG TCATTGAACA ATCATGGATT GTTGTGTTTG 120
GGATTTTTTT TTTTNGGN TTNNTTTTGG GTTTNGNGN GTGTGTGTGT GNCCTGN 177

SEQ ID NO:7429

SEQUENCE LENGTH:388

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08670

SEQUENCE DESCRIPTION:

GATCCAGGTG GTCCAGGAGC CCAGGACAGG CCTTNCCTGT GGGCCCTGGC CAGACAGGGT 60
TACCTGGTGA GGTGCAGAGA GTCCTCTAG TGGCCATTTT GTATGGTAGT TGCTAATGCA 120
GAACAAGTTC TGTCTTGGGC TTAATTGAC TGAAGACTTT AGGGGGAAG AATAGTAAAT 180

EP 0 679 716 A1

GCATGTAAAC AAATGGGGAC ACTCTGTTCA GGAGAATAAT CCGACTGGCA TTTGTGGCAG 240
 TTTTGTAAAT GTAAATGTAT TCATGTGTGT TCTTGTAAT ACGTGTGCT CAGATGTCCT 300
 TTGAAGTGGG AGGGAATCAA TCCGGGGATA ATTTCAAATG GAATAGAGTA TTTTGATATT 360
 GTTCATTCAG AGGGTGATGT GTACACAN 388

SEQ ID NO:7430
 SEQUENCE LENGTH:392
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08671
 SEQUENCE DESCRIPTION:
 GATCCCCACT GTCAATGGGG GATTGTCCCA GCCCCTCTTC CCTTCCCCTC ACCTGGAAGC 60
 TTCTTCAACC AATCCCTTCA CACTCTCTCC CCCATCCCC CAAGATACAC ACTGGACCCT 120
 CTCTTGCTGA ATGTGGGCAT TAATTTTTTG ACTGCAGCTC TGCTTCTCCA GCCCCGCCGT 180
 GGGTGGCAAG CTGTGTTTAT ACCTAAATTT TCTGGAAGGG GACAGTGAAA AGAGGAGTGA 240
 CAGGAGGGAA AGGGGGAGAC AAAACTCCTA CTCTCAACCT CACACCAACA CCTCCCATTA 300
 TCACTCTCTC TGCCCNATT CCTTCAAGAG GAGACCCTTT GGGGACAAGG CCGTTTCTTT 360
 GTTCTGAGC ATAAAGAGGA AAATAAATCT TN 392

SEQ ID NO:7431
 SEQUENCE LENGTH:127
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08673
 SEQUENCE DESCRIPTION:
 GATCTTTGGG AAGGGAGACT AGGGCAGGTG GAGACAGCGC AGAACCCCG TGCTGGGTGG 60
 GAAGCATGAC CACACGGTGG GTGAGCAGCC CCCATGCACT GATGGTAAAT TCCCCTGTGG 120
 ACTCAA 127

SEQ ID NO:7432
 SEQUENCE LENGTH:353
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08674
 SEQUENCE DESCRIPTION:
 GATCTGTCTG GCATGTCAGG AGCCAGAGAT ATTTNATAT CAAAAATTGT CCACAAGTCA 60
 TTTGTGGAAG TGAATGAAGA GGGACAGAG GCGGCAGCTG CCACAGCAGG CATCGCAACT 120
 TTCTGCATGT TGATGCCGA AGAAAAATTC ACTGCCGACC ATCCATTCTT TTTNNNNNTT 180
 CGGCATAATT CCTCAGGTAG CATCCTATTC TTGGGGAGAT TTTCTTCCCC TTAGAAGAAA 240
 GAGACTGTAG CAATACAAAA ATCAAGCTTA GTGCTTTATT ACCTGAGTTT TTAAATAGAG 300
 CCAATATGTC TTTATAANCT TTNCCCANTA AAAACCCACT TGTCCCAAAG AAA 353

SEQ ID NO:7433
 SEQUENCE LENGTH:399
 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08675

SEQUENCE DESCRIPTION:

5 GATCTAATTG GCTTTTATTA GCAATTCATG AGCCCGGCAG CATCCAGTCT AAAAAATAGA 60
 AAGGAGTGCC ACTAAGCTAG ACAGAATGGA TGGGTTTTAT GGCAGAAAAA AGTGGGAAGCA 120
 AGCAAGAACA ATGAACAAAT TAATGGATGG ATTGTTTTAA GATTACATTT CTTTATGGG 180
 AGTAAACAAA ATCTTGTTAG CCTAATGGAA TTTGGCATCA TCTCCTGATT TCTCAGAAGG 240
 10 TCATATCTTG TAAGTGAACC ATTTAGGTTT GGTGACCTGG NACCTTTGGC ATAANGTGAC 300
 TCCATTTTGG GGCCTGCTGC CTTTTTCTTT AAAAANGGAG TGACACCNTA ATTTCCATGG 360
 GNGAGCACA CANNATGCCC TTGTNTTGCA GGACCTNCN 399

SEQ ID NO:7434

15 SEQUENCE LENGTH:230

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08676

SEQUENCE DESCRIPTION:

20 GATCCTGACC ATATTCTCA AGGTCTCCTT TTCCTTCCTC CAGTGTAACA AAATTCCTG 60
 CTGGCGTCCT ATGTAAAGAT AATCGGCCCT TTTGGACCT TTTGTTGGN TCAAGAATTT 120
 GGTGGGAGA AAAGAAAGTG GGTATCAAG GGTGATTGA AATTTCTGC AGCATTAAAG 180
 CTGGCGCTTA ATAAGAATAA GTAATAATAA AGAAATTCT AACATTCAA 230

25 SEQ ID NO:7435

SEQUENCE LENGTH:153

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS08677

SEQUENCE DESCRIPTION:

35 GATCTGAAA ATAGCCCCGT GAAGGCAGAA ATGTATGTGA CTAGAACGAG GCCACATGAA 60
 TAAGCCACTG CCCACTGGCA GGAGTGAAAA CTGAAGCGCT CTTACCTGA AGGACCCCAA 120
 AACCATATAG AATAGAATAA CCAGGAGTTC AAA 153

SEQ ID NO:7436

SEQUENCE LENGTH:138

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08678

SEQUENCE DESCRIPTION:

45 GATCTCACAA GCNCTTTCCC TCTTGGTGCC TCAGTTTCCT GACCTATGAA ACAGAGAAAA 60
 TAAAAGCACT TATTTATTGT TGTTGGAGGC TGCAAANTGT TAGTAGATAT GAGGCATTTG 120
 CAGCTGTGCC ATATTAAA 138

SEQ ID NO:7437

50 SEQUENCE LENGTH:361

SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS08679

SEQUENCE DESCRIPTION:

5 GATCATGGCT AAAGATGACC TCTCTGGTGC TGACATCAAG GCAATCTGTA CAGAAGCTGG 60
 TCTGATGGCC TTAAGAGAAC GTAGAATGAA AGTAACAAAT GAAGACTTCA AAAAATCTAA 120
 AGAAAAATNTT CTTTATAAGA AACAGGAAGG CACCCCTGAG GGGCTGTATC TCTAATGANC 180
 CATGGCTGTC ATCAGGAAAA TGGTTGGGAG ATTTCTCAAT CCCTGAAAGG ATGAGGTTGG 240
 10 GGGAGTTGCC CAGAGGAATC CCTGTTCCTCA CTGATTTTAA TTAGCAAAAC ATCCTGTGTC 300
 TTTTGGAGTA CGNTGTGTAA GTGCCCATTG GGTGGCCTGT TNGGTCACCTG TGCAGCAGTT 360
 N 361

SEQ ID NO:7438

15 SEQUENCE LENGTH:273

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08680

SEQUENCE DESCRIPTION:

20 GATCAAAGTG TAAACACCAT AGGGACCCNT TCTACACAGA GCAGGACTGC ACAGCGTCCT 60
 GTCCACACCC AGCTCAGCAT TTCCACACCA AGCAGCAACA GCAAATNACG ACCACTGCTA 120
 GATGTCTATT CTTGTTGGAG ACATGGGATG ATTATNNNCT GTTCTATTG TGCTTAGTCC 180
 AATTCCTTGC ACATAGTAGG NACCNNGTTC AATTACTATT GNATGAATTA AGAATTGGTT 240
 25 GCCATAAAAA TAAATNNGGT TCATTAAANG TNN 273

SEQ ID NO:7439

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS08681

SEQUENCE DESCRIPTION:

GATCAGAGGA ATGTTATNTT GTTCAGCCTT CTGGTTGGTT CACAGTAGTG ATAGTATTGT 60
 35 AAATACTCAG AGCAATCCTA CAAGAAGAAA ACTACATNAG CCCCCACATG TTCCCCCAA 120
 CATACCCCAT CACAGCAACT GAAAACAAGA GAAATGTTAG TTTTCTGTGG AAA 173

SEQ ID NO:7440

SEQUENCE LENGTH:194

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08682

SEQUENCE DESCRIPTION:

45 GATCTCTTGA ACTCTGGGTT CTCCCAGCCC CTTGTCCTTC CTTCCAGCTG AGCCCTGGCC 60
 ACACTGGGGC TGCCTTTCTC TNACTCTGTC TTCCCCAAGT AAGGGGGCTC TCTNAGTGCA 120
 GGGTCTGATG CTGAGTCCCA CTTAGCTTGG GGTGAGAACC AAGGGGTTTA ATAAATAACC 180
 CTTGAAAACT GAAA 194

50 SEQ ID NO:7441

55

SEQUENCE LENGTH:132

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08683

SEQUENCE DESCRIPTION:

GATCANATGA AGAAAAGGAG AGGTAGATAC AGTCAGTGTC ACTTCAGGAA AGCTATTAA 60
 AAAANCTTGA AATATAATTG AAAGAAGAAA CAACACCAAA NAAGCCTAAN CCTAGCCTCT 120
 GANCANCACT AN 132

SEQ ID NO:7442

SEQUENCE LENGTH:208

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08684

SEQUENCE DESCRIPTION:

GATCAATCTG AGGAGACTTA ATAATGATAT TNATTCTCCC AATTCATGAA TATAGTATAC 60
 CCCTGTATTT ATTTGTNTTC TTGAATTTCT TTTATCATTG TTTTGTAGIT TTCACCATGA 120
 CAGTCTTGCA CATATTTTGT TAAATGTACA GCTGAGAATT TAATTTTTC TGGTGTACAA 180
 TGCTAATAAA ATGGTGCTTT AAAAGAAA 208

SEQ ID NO:7443

SEQUENCE LENGTH:244

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08685

SEQUENCE DESCRIPTION:

GATCCTCTAA GGTCTACAG TTTACTTCTT GTATTCTCCT TTGTAAGTCA TCTCCAAGAC 60
 GATGTCCAAA TCCATCACCA TTAAATTTAA TAGTTTCCTC ACCCACAACA CTTAATATTT 120
 TAAAAAANGA TACTTTTCAT TGTATTATAA TTAATTGATA CATACATATT TGCTCTGTGA 180
 GTTCCTTATT CATCATATTA GTGCCTGACA ATAAATGTGT GCTGGATTGA GCTGAATCTT 240
 TAAA 244

SEQ ID NO:7444

SEQUENCE LENGTH:323

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08686

SEQUENCE DESCRIPTION:

GATCCTGCAA TTCCACTTAT AGGAATTGAC CCACAAGAAA TGAAAGCAGG GACTTGAACC 60
 CATATTTGTA CACCAATATT CATAGCAGCT TATTCACAAG ACCCAAAAGG CAGAAGCAAC 120
 CCAAAATGTT ATCAATGANT GATTGATTGG CTAAGCAAAA TGTGATATGT NCCTAACGAA 180
 GTATCCTTCA GCCTGAAAGA GGAATGANGT ACTCATACAT GTTACAACAC GGACGAACCT 240
 TGANANCTTT ATGCTAAGTG AAATAAGCCA GACATCANCA GATAANTAGT TTATGNTTCC 300
 ACCTACATGN GGTACTGNGA GTN 323

SEQ ID NO:7445

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08687

SEQUENCE DESCRIPTION:

GATCCTCCTG CCTCATCCTC CCAAAATNCT GGGATTACAG GCATAAGCCA CCGTGCCTGG 60
 CCTCTTTAAT AATNTTNAAT ATACCCATAA GGCTTGTGAA TATACAAGTC TACTGATAAA 120
 TTATGTATTG TCTGGGAATT TGATAGTNAT TGTTTATAGT AACTGGNNTT TACGCTGTGG 180
 TAGNCNGGCT GTGACACTAG TGTTGCACAG GTGTAATTGG TCATCCTATG CCTTCACCAG 240
 AATAACTTGG GAGTGGTGCC AGAANCTAGA GTCTACAATT CTCACTGTTT AGAGAGTGTT 300
 ANTGNCATAC TGTGTATN 318

SEQ ID NO:7446

SEQUENCE LENGTH:317

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08688

SEQUENCE DESCRIPTION:

GATCAAGATA GTAGTATTAT TACACAAGAA ACTTGGTCTG CAGTCTGGAA GCTTGTCTGC 60
 TCTATAGAAA TGAAAATGCA GCATGAAGTT GACATTGTGG AAATGAAAGT AATTGGGTAT 120
 TAGAAATCTG AAAGTACTGT CATCTAANAG CAATTGTGAT TTTNTTGTA TGGTGTGCA 180
 CTGTTGTNCG GTGTCTAGNG TTANNGAATA CATGTNANCT TTCATGGTAT TTNGCCTTTC 240
 TTAANTTTTT TTAATAATTA ANCTTTCTAA CCTATGTATT CAACTTCTGT ATTTATATNT 300
 NATCAGTGGT TCATGTN 317

SEQ ID NO:7447

SEQUENCE LENGTH:284

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08689

SEQUENCE DESCRIPTION:

GATCATCAAA GAAAGACAGG GTGACCTGAG GATTATATGT TCAGCCAAGC TCCCCTCCAA 60
 GCATAAAGGC TACAGATAAC CATTTGTACA TATTCAGGAA CTGGAGTAAT GTGATTCCAA 120
 TGAGCCTGTC TTGAAGGAAC TCCTAGGGGA TTAACCTCAG CTAACCTGGG AAGTTATCTG 180
 GAGAAACCAC AACAGAAGGA CTGGTTGTAG TTCGAGGCTG TAGTCAGCTA TGATTGCACC 240
 ACTGCACTCT AGCCTGGATA ATAGAGCAAG ACTTTGTCTC CAAA 284

SEQ ID NO:7448

SEQUENCE LENGTH:324

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08690

SEQUENCE DESCRIPTION:

GATCCCTAGT ATAACACATT CAGTGTTCCT CTTTCAGTCT TACTACTTTN ACCGCGATGA 60

EP 0 679 716 A1

5 TGTGGCTTTN AAGAACTTTG CCAAATACCT TCTTCACCAA TCTCATGAGG AGAGGGAACA 120
 TGCTGAGAAA CTGATGAAGC TGCAGANCCA ACGAGGTGGC CGANTCTNCC TTCAGGATAT 180
 CAGGAAACCA GACTGTGATG ACTGGGAGAG CGGGCTGANT GCAATGGANG TGTGCATTAC 240
 ATTTGGAAAA AANTGTGAAT CAGTCACTAC TGGANCTGCA CAANCTNGCC ACTGNCAAAA 300
 TTGACCCCN TTTGTNTGAC TTCN 324

10 SEQ ID NO:7449
 SEQUENCE LENGTH:324
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08691
 SEQUENCE DESCRIPTION:
 15 GATCACCTGA GGTGAGGGT TCGAGACCAG CCTGGCCAAC ACGGTGAANC CCATCTCTAC 60
 TAAAAATGCA AAAATTAGCC GGGCGTGGCG GCACATGCCT GTAATCCCAG CTACATGGGA 120
 GGCTGAGGTG GGAGAATTGC TTGAACCCAG GAGGCGGAGG CAGAGGCTNC AGTGACCCAA 180
 GATTGTGCCA CTGCACTCCA CCCTGGGCAA CAGAGCAAGA CCCCATCTCA AAAATAAATA 240
 20 ANTATATATA NAAAAATAAA NGCTATTTCT NGNNNTTTC ACTATAAAGT TTTGCTTTAT 300
 TAAAAAGCTA NTANGCAGCT ATTN 324

25 SEQ ID NO:7450
 SEQUENCE LENGTH:326
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08692
 SEQUENCE DESCRIPTION:
 30 GATCTTAAAC TCAGTAAGCC ACTATCTGCA ATTTTGTACA TTATATAGTA TTTTGAAGAT 60
 ATGGAACCTT ATGAAAAAAA AATAGCAAAT NAGTTCCTTT TCCCCCAGAG GGGAAAGTTA 120
 TGTNCTGCAA ATAGTGTGTG TCTTATTTA CTGTTGANCA GCAATNGCNN NTTATTTTTT 180
 NATTGCCCTAG AACTTCAACA TGTGTATAG GAATCCTGTA GTGCCACTAG TTAAATGCCG 240
 35 AATTCTCATC TGGATGTTAC CATCAAACAT CAGTACACTT GTCATTTAC ATGTGTTTAA 300
 TGTGACAGTT TTTCAGTACT GTNTNN 326

40 SEQ ID NO:7451
 SEQUENCE LENGTH:38
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08693
 SEQUENCE DESCRIPTION:
 45 GATCTNGTGG GCATAAATAA AGGTGTCATA AAGACAAA 38

50 SEQ ID NO:7452
 SEQUENCE LENGTH:83
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08694

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EP 0 679 716 A1

SEQUENCE DESCRIPTION:

GATCTTAAGA AGTGCTTTGA AGGGTTAAGA ATCAGGGGTC CAAGAGAGAC CCCAGTCCCT 60
CAATAAAGCC ACAAGAGCCC AAA 83

SEQ ID NO:7453

SEQUENCE LENGTH:351

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08695

SEQUENCE DESCRIPTION:

GATCTACACT CCAGGCCGGA AAGAGCAAGG GGAGCCCATG ACCCCTCGCC GCACGCCAGC 60
CCGCTTTGGT CTGTAAGCTG AGGCTGTCCC CAGAGAGGAT GGCAGCAGGT ATTGGGTCCT 120
CAGCCTTCTG GCGGGAGCCT GAGGCTGCGG ACAAAGCCCT TTAATCTAAG GACTTTAATC 180
TGTNCATATN ACGGCCCCC AGGGCAGTTC CTGCTGGACC AGACTCTNTG GCAGAGGAGG 240
TGGAGTTCTT CCATGCAGGA GCACGNCATG GCGGGAGCGG GGCTGCAGAG TATCCGNGGT 300
GCTGCCGNGG CACGGNGAGG TNGCTNGACC CATCGCATCT AAAANTAGGA N 351

SEQ ID NO:7454

SEQUENCE LENGTH:55

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08696

SEQUENCE DESCRIPTION:

GATCTAAGGT TTTAAAGAA TTTGGTTGA AAATATACAG TTGTCTTGAA GGAAA 55

SEQ ID NO:7455

SEQUENCE LENGTH:61

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08697

SEQUENCE DESCRIPTION:

GATCTGTTCT GTTTCACCAT GTAACACACA ATACATGCAT GCACTGTATT AGTGTTAGAA 60
A 61

SEQ ID NO:7456

SEQUENCE LENGTH:273

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08698

SEQUENCE DESCRIPTION:

GATCCCATTT TCCTCTGACG TCCACCTCCT ACCCCATAGG AGTTAGAAGT TAGGGTTTAG 60
GCATCATTTT GAGAATGCTG ACACTTTTTC AGGGCTGTGA TTGAGTGAGG GCATGGGTAA 120
AAATATTCTT TTTAAAGAAG GATGAACAAT TATATTTATA TTTCAGGTTA TATCCAATAG 180
TAGAGTTGGC TTTTNTTTT TTTTGGNCCA NGGGGGGGGG TTTTTCCTCC NTGCCCCCCT 240
GGGGTTTTTN NANTCCCNCT NNNAAAAAAN ANN 273

EP 0 679 716 A1

SEQ ID NO:7457
 SEQUENCE LENGTH:363
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08699
 SEQUENCE DESCRIPTION:
 10 GATCTTAACC AGTCGAGTGG AGTGACATT GTCTGAATAC AGGNTGCACA ATGTTGTCAA 60
 TCCTGGAAAT GGTCTTTCTT TTTTGTAAGA TATGTGAATG AAGTGTGGT GTCCTCACCA 120
 AGAGGTGGCA CCTAAGGGTT CTGAGGAAAT AAATGTATAG ACCCTTATGT ACAGACCTGT 180
 GTATAACAA CTNTTGATA TACATATAGG ATAGCTTTT TGTACTATAC AGCTGGTACC 240
 ATAANAGTAG CTGGATANTT TGGTTAGGNC NGGTTGTCCA CCAGTTTGG NNTTTTTTT 300
 15 CACTTTGNTN CATTTTGGGG GATTTTCCTT GTGGGTGNN TAAAANTTG CATATGCTTA 360
 NGN 363

SEQ ID NO:7458
 SEQUENCE LENGTH:173
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08700
 SEQUENCE DESCRIPTION:
 25 GATCCGTATT TCCAATAAAA TCCCCTACCT GAATATTCAA GTTAAACATG TCCAGAATAC 60
 TTACTGATTT TATTGTTAAT AGCCACTATT CTGTTGTCTG GAATTA AAC CTGTATAACT 120
 AATTGCATC CCTTTATCTT CTTAGTCAAT AAAACCTACA ATCCTCTTC AAA 173

SEQ ID NO:7459
 SEQUENCE LENGTH:349
 30 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08701
 SEQUENCE DESCRIPTION:
 35 GATCACCCCC CAACTTGCCC TTCAAGTTCT ACTTCAGTTT GATAAGGCTA TAAATGCAGC 60
 ACTGGCTCAG AGGGTCAGGA ACAGAGTCAA TTTCAGGGGC TCTCTAAATA CGTACAGATT 120
 CTGCGATAAT GTGTGGACTT TTGTACTGAA TGATGTTGAA TTCAGAGAGG TGACAGAACT 180
 TATTAAAGTG GATAAAGTGA AAATTGTAGC CTGTGATGGT AAAAATACTG GCTCCAATAC 240
 40 TACAGANTGG ATAGAAAANN TATGACTNNT TTACACCATC TTCCTGTTAT TCATTGCTTT 300
 TTGAAGAGNA GCATAGANGA GACTTTNTN TTTTATNCTN GGAATTTN 349

SEQ ID NO:7460
 SEQUENCE LENGTH:395
 45 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08702
 SEQUENCE DESCRIPTION:
 50 GATCTGGAGT AAGGAATATT GTCATCACCT GGATTTTGAG AAAGAAAAAT AACTTCTCTG 60

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EP 0 679 716 A1

5 CAAGATTTC TAATTGAGAG AATTCCTGAG TTGATAGCTC TAAAGGCAGA TATGCTGTAT 120
 TTGCCTACTT TAACCCATTT TTCAACCTGT TTGTTTTTA ANNGGNTTCA CTAAGGGTTG 180
 ATATGTACCA TTGTATGGGG CAATTTTAAG TCAGCTAAGG CAATAACCTT ATGCATGAGC 240
 ATTTCCCAGA CTTTCATGAA GCTGTTGAGG TCCTAGGCAA TTAATGCAGC AGTTGCGATA 300
 AATAAANGNC NTCTCACCTA AGNNTCCNTT TTCTTCNATA ACCATGGTTC CTGGNCATGN 360
 TTGGGAGGCT NCTCGGCTTT AGGGNAANGG GGTTN 395

10 SEQ ID NO:7461
 SEQUENCE LENGTH:347
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08703

15 SEQUENCE DESCRIPTION:
 GATCGCCTCC CAACAGCAGA TTTCGACATT ACCTGAGAGT CTTGATTTTA GGCTTGTTTT 60
 TTGTAAACCC ATGTGTTTGT AGAGATTTTA GCGCTCTTCG GATATCTTCT CACCTATGTT 120
 CCCTGGCTAA GAAGTCAGAG GTAGCCAATG TTTCCTTAAA TTCATTTTA AACTTACCAT 180
 TGGTGACATAT GTTCCAGATG GCAGATGCTG TCAATAATCT CACCATTGAT GACCTTTGTG 240
 20 TATGTAGTTC TTGCATCCTA TACTGGATAA GCCTGTTTTA ACCTGCTATG ATGGGTGCTT 300
 TCCATTGCTT CNTAATCTTC ATGGAGGGTT GCCATGCTTT TTGCCNN 347

25 SEQ ID NO:7462
 SEQUENCE LENGTH:274
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08704

30 SEQUENCE DESCRIPTION:
 GATCGGNATG TAATAACAGC ATTAACCTCAC AGACCTTGGA GCCTAAGCCA TACAGGAGAT 60
 GGGAACCCAC GCTATGATAC TTNCTGGAAA CATTTTATAT TTNTAATGAT GGACATTTTN 120
 CTCGATTGGA GCATGCATAA TANTTGTGG TACCTGTGTG GAATTCANC TTTCCTCATG 180
 CAAAAGGATT TTNTATCCCC GGCCTACTTG ANGAAGTGGT CAGCTAANGG NNTCCAGGTT 240
 35 GTTGGTTGGA CTGTAATACC NTTGATGAN AAGN 274

40 SEQ ID NO:7463
 SEQUENCE LENGTH:69
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08705

45 SEQUENCE DESCRIPTION:
 GATCAGAAAC TTGCATATGC GTGAAAAAG GAAGAGCAGC AGAAAAATAA TAAATGACAG 60
 GAAAGCAAA 69

50 SEQ ID NO:7464
 SEQUENCE LENGTH:199
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08706

SEQUENCE DESCRIPTION:

5 GATCTATAGC AATGTGATTT TATTTTAA AAGAAAAGCA GTGTGTTTC TTTTTTGTT 60
GTTTTCTTTT GCTTAAGCAC TTCATCAATT GCTTTATTCT GTATCTGCGA AGTNATCTGC 120
AATCTCTNTT GNNCTTTNTA AAATTTGGAT NTGTTATAAA AATTGNCCAN ATTGGAAGGT 180
GTTTCAGTTA AAAAAANAN 199

SEQ ID NO:7465

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SEQUENCE LENGTH:320

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08707

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SEQUENCE DESCRIPTION:

20 GATCACAAC TTGGAGGGCT GTATGGTTTT TCATTTAAGA AAGTAATTGT CGTGCAATAA 60
AGTTATTAAG CCTTGTTTAA GAATATTTTT ATTTGTACA TCAAGTGGAG TTGGAATAG 120
TTGGTTTCAT AATGATATAT TTCAAAAGTA TCTTGAAGTC ATCTGGTTCA TTCACTACC 180
CAGGGCACAC ATTCTCCTG TAACATCCCA AGTGGTTATT AAGCAAAACA GATTTATTTT 240
ATTTAGAAAA NNTTAGTATG ACTGATTGTC TATTGATTTT GACTTCTAT AAATNAATGT 300
GGATACATGC TATATTCAAA 320

SEQ ID NO:7466

25

SEQUENCE LENGTH:311

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08708

SEQUENCE DESCRIPTION:

30 GATCACTTAC AGGTGCCCTC TCCCCAACA GCCAGAAAGG GAAAACAGAG CCGAGTTTGG 60
CCACTGGGGC CTCCAGAGCC CTCTGGAAGA AGGGGCTGAC TGATAGGAAG GCACTCCTCT 120
GGTGTTCCT GGCAGAGCC CTGTGGGCTC AGAAGACTAC CTAGACCCCA TGAAGAAAGA 180
CTCTAGACAC ATTTTATAA TCAGAAACTT ACAGTTTATT GTGAATAAAC CACTCAACTT 240
TGAGCAGATG ACATTGCCCTC ANTTAACTTT TTCGTTGAAG AAAACAAAG ATATATGGTA 300
AATTCATGAA A 311

SEQ ID NO:7467

40

SEQUENCE LENGTH:434

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08709

SEQUENCE DESCRIPTION:

45 GATCATCAAC AAAGACTTGT TAGAAAGGTC TAGTCTTAGC ACTTGGCAGT TAATCTAGGG 60
AAGATGAATT AAATGGGTAG ATAGTGATGC ATACCTGTAT TCACTGATGT ATGTTTAAGG 120
GATTTGGGGG GGATACCTCA GTTCATGTGG AAGGGACAGT CTCGGTGTGT CCCATGAATA 180
ACCTTGGAAC TGCAACAAAT GGTGTGCT CAGAAAAAGT CTTTCATGGT GACAGGAAGA 240
CAGTTTCCTT GGAGCTGGCC ATGAAGGCCT TAGAAGCCAT TTCTGGTGTG TGGTGGGTAG 300
50 CAGGCATAGA GATGATGTNC CGAGGTCCCA GTGAACACCA GTAGCCAAAN GAATTGTACT 360

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EP 0 679 716 A1

ACCTTNTATC CATTANTTNG GGGAAAGGTCC TTCCCTNGGG NAACCAATGN CCTTTTTTGN 420
TGGGCCAAAA ITTN 434

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SEQ ID NO:7468
SEQUENCE LENGTH:414
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08710

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SEQUENCE DESCRIPTION:
GATCGAGCTG TGGACTTATA CCTGTGTGCN TGGTTTACAG GTCCAGTGGT TCTTCAGCCC 60
ATGACAGATG AGAAGGGGCT ATATTGAAGG GCAAAGAGGA ACTGTTGTTT GAATTTTCCT 120
GAGAGCCTGG CTTAGTGCTG GGCCTTCTCT TAAACCTCAT TACAATNAGG TTAGTACTTT 180
15 TAGTCCCTGT TTTACAGGGG TTAGAATAGA CTGTTAAGGG GCAACTGAGA AAGAACAGAG 240
AAGTGACAGC TAGGGGTGA GAGGGGCCAG AAAAACATGA NTGCAGGCAG ATTCGTGAA 300
ATCTGCCACC ACTTTATANC CAGATGGTTC CTTTCACAAC CCTGGGTCAA AAAGAGAATA 360
NTTGGCCTA TNATGTTANA AGGAANGCAG GNNGGTGGGT TNAANTAAAA NTTN 414

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SEQ ID NO:7469
SEQUENCE LENGTH:214
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08711
SEQUENCE DESCRIPTION:
GATCAGTATT TTTAATCAAC ACCTGAGAAC TGTTACACCT TTNATTTTGT CTTNAGGAA 60
ATCCCTGTCT TTCCATTTT TCATGTAAAT NTTGCACAGT TACTTGTTCA TATGTAAATA 120
TTTACTTTC AGAAATNAAG TTTTAATTG CTATTGTTTT ATATAGGATT GAAAGAAAAT 180
30 NAACTCCTTT ATTAAAAACA AATTTATCTG TAAA 214

30

SEQ ID NO:7470
SEQUENCE LENGTH:418
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08712

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SEQUENCE DESCRIPTION:
GATCCATTTC CTCAGGACTC AGTTAGGTGT GCCCAGACGC ACTGACAAAA TNACGTGACT 60
40 TCAGGGAAGC CTGGACACCC GAGGCACCTG GACCAGCTAT GGGTAGTTCT GTGGGTGGAA 120
CACATTCTGT GTAAGAGCCC CACTGAGGGC TCTGCAGCGG AGTGACAGCA ACCCCAGAGA 180
TGAGGCACCA GAGAGTGCCA CTGCATGAGA CACCTGTGAC CATTGGAAGT CTGAAATGCG 240
GGGGGGGAGT TTCATTTTTA AGTGAAGACC AAAAGCCCTT TAAAAATAAT AGTTTTTTAT 300
45 CATTTTATAG TAATCAGCGG NNTCTCTTT ACTAATATAC TCATTCCTNT TCGGTATGAT 360
TGGNCTTTTG GAGCTACATG CTCCNACAAN NCTGTATCTT AATTGGTTAA CTTGTTAN 418

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SEQ ID NO:7471
SEQUENCE LENGTH:422
SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS08713

SEQUENCE DESCRIPTION:

5 GATCTTGGCG TTGTGGTAGA TTGCCTTCCT GTGCTCACCA ATTGTTTACA GGAAGAAAAA 60
 CAATATATCT CACTTGGCTG CTGTGTTGAC TTGTTCCTC TAGTAAAGTC ACTACTTAAA 120
 AGCAAAATTN AAGAATATGT TATAGTTGGT TAAACTGGC TTCAAGCAGT CATTAAAAAGG 180
 TGGTGGTCAG AACTATCATC CAAAACAGGA ATTATAAATG ATGGAAATAT TCAAATTTTA 240
 10 AAACAACAAT TAAGTGGATT ATGGGTACAG GANAACCATC TTACTTTGGT TCCAGGATAT 300
 ACTGGTAATN TAGCTANGGG TGTAGNTGCT TTTTTCTTT CCAGTTCNT TGNGGGGATT 360
 CCTCTNCTG NGGGGCATT TGGTTTTCA ANACCNTCCC NTGGACCTGT TTTANTTTTC 420
 CN 422

15 SEQ ID NO:7472

SEQUENCE LENGTH:359

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08714

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SEQUENCE DESCRIPTION:

GATCGCCTTA ACTACTGTAA ACATAGCCTA TTTTGTGCT TAANATACTG AATGGAAAAC 60
 TCCATTGTGT GTTGCTGGAC TGTTTGGAA ATATNNGTT AAATGTGTGT TAATTGGCT 120
 GTAATGGCAT TTAAGCAAA CAAACAAACA AAAAAAGCTG TGAAAATGGC CTGGAGCAT 180
 25 TATCTTTAGT TACTTGAAGA GTTCTAGTT TTTTAAAT ACAGTTTATG TTAATAAT 240
 TTTANTTAAT TTAGAGANGC CAATCATTGT CTGNGGGGAA ACCGGCCTN CNNTTGGGN 300
 TTNCNTTTN GTGGGCCATT GGTGAGGTGA GTGCCTNTN CCCGTTTCNT TNGGTTTCN 359

30 SEQ ID NO:7473

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08715

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SEQUENCE DESCRIPTION:

GATCTTTAAA GTTATTTTAA GGCACATTG TGACAAACAA CTAATTGATA TTGAAATCTC 60
 CNTCAGCCAT TAGAAGCTAT CAAATAAAGT AGGTGTAAAA ACAA 105

40 SEQ ID NO:7474

SEQUENCE LENGTH:104

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08716

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SEQUENCE DESCRIPTION:

GATCAAGTAT GCTAATNTTA AGCAAATGTA AAAACTCATA AAACAGGTAA ACAGTGGGGT 60
 GATTTCATT GCCATAATC ACATAAGACG AATTTAATC TAAA 104

50 SEQ ID NO:7475

SEQUENCE LENGTH:340

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08717

SEQUENCE DESCRIPTION:

GATCCTTGTT TTTAAGTTAC TTTCGATGTG TATACGAACA AAGACCAGGG AGAAACAGAA 60
 CTTTAAAAAC TATAGGCTGC TGTNTTGGGG CCAGAAATAA GGTGACAAAGT AAAGACTTTT 120
 ATAATACCTA CCTTCAAANT TAGAATCAGC AGCACTGTAC AANTAAAGTG CCTGTGCTTC 180
 ACCTCACACC AAGANTCTTG CCTAACCCTG TCAGAGTGCC TAATATCCTG TGGTGATTAT 240
 GTATAATGAA ACAGTCTTAT TTTGTGCTCT TTGGTTTCTT AAAAGGAAAG CTTGTTTTCT 300
 TTCCTTTATC ATTACAGAGG TATGTCTTGG GTTCTTANTN 340

SEQ ID NO:7476

SEQUENCE LENGTH:189

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08718

SEQUENCE DESCRIPTION:

GATCTGTGAA AAAATACAGT ATTGTAAATC CAAGACTCTG ATACTGAATT TAAGTTATTT 60
 CACAGTTGTA GCTACACAGT AAGTAGCTTG GTAGATAGTT ATTGANTGTA TTTATGTAGT 120
 GTATTAAGAA GCTTATATTA CTACAAAAAA CTTATTTTAA TATATNTNNN TATNTNNGTN 180
 TTATTTATN 189

SEQ ID NO:7477

SEQUENCE LENGTH:342

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08719

SEQUENCE DESCRIPTION:

GATCTGTGTG AAAAAATNAT TTCTTAGCTA TAATCAGCAC ATCACTTAGT TCAAACAAAA 60
 TTCCCGACGA AATGTTAGAT AGTAGGTATA TCAGTCACCT GGGGAGTTTT CTTCATAATA 120
 TGCATATTCA TCTNGTAATG CATACATAGT TATCATCCTC CTTCTCAACC CATCTCCCTA 180
 ACCCCACATG CTTGCCAGTT CTTGAAGGGA TAAAGTGATT GTAATAGTGT TTNACTNCTC 240
 TCTGTTCAAT TTANTGTGAT ATAATTCTAG TNTAANNNTA TTTNGGACAG TNGCTTAACC 300
 ATGGTCATAA GAGGATTGT ACTATAGATT ATCTTCNGGG GN 342

SEQ ID NO:7478

SEQUENCE LENGTH:54

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08720

SEQUENCE DESCRIPTION:

GATCTTTGTC AACCAAAACA TGTATTTTCA ATAAATTGAG ATTCCAGTT CAAA 54

SEQ ID NO:7479

SEQUENCE LENGTH:79

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08721

SEQUENCE DESCRIPTION:

GATCTGTTTC CCTTTCACAT TTTCCCAATT TTGTGAGCAT TTCCTTACTT CCTAACATGA 60
CAAAATATTT CAAACTAAA 79

SEQ ID NO:7480

SEQUENCE LENGTH:281

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08722

SEQUENCE DESCRIPTION:

GATCCCCCTG GCCTGGAGGC AGCAGCTGTA TAACTCTAC GAGGCCGACT TTGTTCTCTT 60
CGGTACCCC AAGCCCGAAA ACCTCCTCCG AGACTGAAAG CTTTCGCGTT GCTTTTCTC 120
GCGTGCCTGG AACCTGACGC ACGCGCACTC CAGTTTTTTT ATGACCTACG ATTTTGCAAT 180
CTGGGCTTCT TGTTCACTCC ACTGCCTCTA TCCATTGAGT ACTGTATCGA TATTGTTTTT 240
TAAGATTAAT ATATTTCAAG TATTTAATAC GAAATGTGAA A 281

SEQ ID NO:7481

SEQUENCE LENGTH:326

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08723

SEQUENCE DESCRIPTION:

GATCTGAGTC CAGGCAGTGT TTTAAGNGA TATGAGTTAG TAGTGAAAAT TGAAAAGGAT 60
ATTTTACAAC CAAATACAGG TTTNNTGGTG TTTTTGAAA AATCCAAAGA TTTGGCAACT 120
CGACCTAGGC TCCGACAAAA CCACAGTTGT TGGGAGCTGA GCCGGAGCCA GAACTCACAA 180
TTCACCGTGG CCCGGCCACC CATCCCCACC ACTCCCGTGG GCTCCTGACC AAGCACCACG 240
TGCCATTCAT CATGGCTTGC TTTTAGTGTT TTTGGTGTTT ATCTTAAAGT ACAGCGAAGA 300
AGAAAGGGAA GTATTTNTTA GACCCN 326

SEQ ID NO:7482

SEQUENCE LENGTH:331

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08724

SEQUENCE DESCRIPTION:

GATCTGGATT TGGTAAAGGT TGACTTAATT GAACTCTCTG AAAAATGCTG TAGTGACTTT 60
AATTTGCACT CAGAATTAGA GCGCTCATTT TTGTCAGANC CATCATCTCC AGGAAGAACC 120
AAGACTACTA AAGGATTCAA ACTTGGGAAG CACAAGCATG AGACCTTTAT AACGTCAAGT 180
GGAAAATCTG ANTACATTGA NCCTGCCAAA AGAGCTCATG TTGTGCCACC ACCAAGAGGA 240
AGGGGCAGGG GTGGNTTGG ACAGGGTATA CGACCTCATG ATATTTTNG TCAGNGNAAC 300
CAGNNACAN GTNGACCACC ATCTNTGCNT N 331

EP 0 679 716 A1

SEQ ID NO:7483

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08725

SEQUENCE DESCRIPTION:

GATCGGAGTC GCCAGGCAGT ATATCCAGTT AGAACTTCCG GGCTTTTGCA TTAGCTTGTC 60
 TGATGCTCAT GCCCCACTCA GAGAAAAGAC ATCAGCAAAT TAAGAATTTT CTGGGTTCCCT 120
 GTGACCCTCA GGTATTTTA AAGCAATTGG AAGAGCATAT GAACACGGGC CAGCTAGCAG 180
 GATTTTCACA TCAAATTAGA AGTCTGATT TGANTAATAT CATCAATAAG AAGGAGTTTG 240
 GGATTTTGGC AAAGACCAAA TACTTTCAAA TGTGGAAGAT GCATGCGATG AATACCAACA 300
 ATATCACTGA GCTAGTGAAC TATTGGCAA ATGACTTN 338

SEQ ID NO:7484

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08726

SEQUENCE DESCRIPTION:

GATCCGCCCT CCTAGCCTC CCAATCCTCT CTAAAAAAG TGATAGCTCA GAAATATTTG 60
 TAAAAGCAAG GTTTTTATTT CATTTTGGCT CTGTCATTTT CAGAGGCAAA GAAGTTGGCC 120
 TGTAATAAG AGTGCTAGAG CTCTTACGCC CCTCCCCTTC TTCCCACTT CCTACTTCCT 180
 AGCCCTTTTA TCAACTCCTA GAATAGTTAA AGAGAGACAC ATCTAGATGG GATGAAAGGT 240
 GCCCTAAGCA GGAGAACCCTG ANCAAAAAGGC TAGAGGCATG GGCCAGGTAA AATTTGGGGC 300
 CTAGAGTGAA GNCTGTGCTG TCGTTAAGNG CTTTCGAN 338

SEQ ID NO:7485

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08727

SEQUENCE DESCRIPTION:

GATCACAGTG ACAAAGAAGA TGAACAGCCT CAAGTGGTGG TTTTAAAAA GGGAGACCTG 60
 TCAGTTGAAG AAGTCATGAA AATTAAAGCA GAAATAAAGG CTGCCAAAGC AGATGAAGAA 120
 CCAACTCCAG CCGATGGAAG ANTCATATAT CGAAAACCAG TCAAGCATCC CTCAGATGAA 180
 AANTATTCAG GTTTAACAGC AAGCTCAAAA AAGAAGAAGC CAAATGAAGA TGAAGTAAAT 240
 CAGGACTCGG TCAAAAAGAN CTCACAAAAN CAANTTAANA TTAGTAGCCT CCTTTCTTTT 300
 GACAACGAGG ATGAAAATGA GTAAGTGTA TTNTTTTN 338

SEQ ID NO:7486

SEQUENCE LENGTH:302

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08728

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

5 GATCTTAATT CATATCTTGA AGATAAAGTC TACCTTACAG GGTATAACTT TACATTAGCA 60
 GATACTAT TGTACTATGG ACTTCATCGC TTTATAGTGA GTATTTGTAT AGTTACTGGC 120
 TTTTCTCTG TAGTGTGAG AATGATTTT AAAACTTTGT CTTGAATTA AATCAGATTT 180
 GCATTTTGG CTTATCAAAT AAAAATGAAA GGAATTCCTT TATACAAGGT GCCTTTTGT 240
 ACAAACCTGC ACGTTGTGAC ATGTACCCTA AACTTAAAG TATAATANTA ATAANNTTNA 300
 AA 302

10 SEQ ID NO:7487
 SEQUENCE LENGTH:334
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08729

15 SEQUENCE DESCRIPTION:
 GATCAAAAGT GAATAATGTA TGAATGAGAG CTGTAAGAAG GATTTTNATT TTGTTATAAT 60
 TTAGTTACCA TTTTCAGTGT TATTTCAAAG GTTCTTTGAA GAATTTTGGG GCAGGGCATC 120
 AGATTAGAGT TTTAAATTT GAGTATTTG GATATCAGTG TTCCTCATGA AGATATACAT 180
 GGATATNCAA TTTTGATGGC TTCCAGATTT GTAAGATTTT ATGTTGTATA TACCATTCTA 240
 20 TTAAGAAACA TGTCCACTGT GCTTTCAAAC ATAGATAAAG CATGATAAAG ATTATNATTT 300
 AAGATATACT TGTATTTATA CCTCAGATAT TCTN 334

25 SEQ ID NO:7488
 SEQUENCE LENGTH:337
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08730

30 SEQUENCE DESCRIPTION:
 GATCAGCTGG ATACTCNTGT NACCAGAATT CCAAAGGTTG TGATAGAATC AAAGATGGAT 60
 ACAAAGTGAA CTCACACATA GCTAAGCTGC AAGAGTTATG GAAAACCTCC CAAATCAAA 120
 CAATCCACCT CTCTAANTCA NTGATGGAGG CGTCCTTTT CAAGCATCCA GACCTCACCA 180
 CAGGCCAGAN GCGTTACCNG TGCAGCATTG CTAAANTCTA TAATGCAAAC TATCTGANGA 240
 35 TGTTAATGNA GNGGCAGTAC ATGCACGTNC TTCAGCACAG CTCACAAAAG CCAGGTGTCC 300
 TCACTCATCA CAGNNGCCGC CTNGCTCCC GTTTATN 337

40 SEQ ID NO:7489
 SEQUENCE LENGTH:292
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08731

45 SEQUENCE DESCRIPTION:
 GATCATTGAG AACACTGNAC ACAGCAGCTT CAGGCAAAGG GAAAAGAACT AAATGAATTC 60
 CGGGAAAAGC ACAACATTCG TCTCATGGGA GAAGATGAGA AGCCAGCAGC CAAGGAAAAC 120
 TCAGAAGGGG CTGGGGCTAA GGCCAGCTCA GCTGGAGTGT TGGTCTCCTA GGGACCAAGG 180
 CCTTTGCATT TTTTCTACC CTGACTCCCA CTTCTAATTT CTCTTTATTG TTATTATTAT 240
 50 TATTTTCTCT GCTATTGTAA TATTTTTTTG TTAATTAAT GTTTTGGTTA AA 292

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SEQ ID NO:7490

SEQUENCE LENGTH:140

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08732

SEQUENCE DESCRIPTION:

GATCAACTGC AACACAGGCAC GAAGGAACTT TTTGGGTGA TGAAAAATAA AACTTTGGAT 60
 GTCGAACATA TTTTGAAACT AGATTGCAAT NATAGTTGCA TAACTNTATA CATTTATTAA 120
 AATCGTGGAA CACTTACAAA 140

SEQ ID NO:7491

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08733

SEQUENCE DESCRIPTION:

GATCCGGTCT CCCCCGCCCC CCGGCTGCAG GGTGGATAAA GAGGAAGATG TGAAGAGGAG 60
 GCAGACAAAA GCAAAAGGAC ATCCTGTCAG TAGGAAACGG CCGAATTACA ATTCAGATGT 120
 GCAATGTTTG CATAAGTAA CTGACACGGG ACTATTATAT GCAGAANTCC ACGTGCAAAA 180
 NTTTTAGAC ACACCTTTGT ACCTCTNCCT AATCCTGCCC TGGTAAAATG GCGGTTATCA 240
 GTGCAAGTTT GAGGATACAT ACAGCTGGAA ANTGGNCATC ACCAAACTGG ACTCCAGAAA 300
 GTNCTGTCTA GGATTTTATT AGACTGCTTT TTTNCAN 337

SEQ ID NO:7492

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08734

SEQUENCE DESCRIPTION:

GATCTATTTC TCAGTAATNA GTGTGTGCCT CATGCCTGTT TCAATATTGG GTTTTGGAGA 60
 CANTATTGTA CCAGGCCTGT TGATTGCATA CTGTAGAAGA TTTGATGTTT AGACTGGTTC 120
 TCCTTACATA TACTATGTTT CGTCTACAGT TGCCTATGCT ATTGGCATGA TACTTACATT 180
 TNTNGTCCTG GTGCTGATGA AAAAGGGGCA ACCTGCTCTC CTCTATTTAG TACCTTGCAC 240
 ACTTATTACT GCCTCANTTG TTGCCTGGAG ACGTAAGGAA ATGNAAAAGT TCTGGAAAGG 300
 TAACAGCTAT CAGATGATGG ACCATTTGGN TTGTCAN 337

SEQ ID NO:7493

SEQUENCE LENGTH:75

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08735

SEQUENCE DESCRIPTION:

GATCGGCCTG CGAAAGGATA CTGTGAAATN ACTAATTAAC TAATAAACCT GTCTCAAGTT 60
 GAGGATTTGA AGAAA 75

EP 0 679 716 A1

SEQ ID NO:7494

SEQUENCE LENGTH:343

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08736

SEQUENCE DESCRIPTION:

GATCTGGGGT ACTTGGCAGG GCTTCCCTCA GACAATACCT TAGAGAGTCA CAACCATTCC 60
AGGGTCTGGT CTACCTCACA TCCTCATTCA TTGTTGGATG TGCTTAGCAT AGCCAGAGGT 120
GTCTTCTCTT TGACAGACTT GCTATCTGAA AAATAAACCA GGACACCAAT TTTTTCAGCAT 180
ATAATAGAGG ACATCCCTTC CCAGCTATAG GAATGCTTGG TATCCTTCCA AGAACTCCTC 240
CTTTAGAGGA GCCTCACAGA AGTTGCTGGG TAAACATGGG CTCAAACCAA GGAAAGTCAG 300
GTAACCTGGG TGGNGGTAT CCCACCACCA ATGGCAGGGG GTN 343

SEQ ID NO:7495

SEQUENCE LENGTH:95

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08737

SEQUENCE DESCRIPTION:

GATCAATATT GTAAAGAATT TATTCTGATA AATNAGAAAC TGGATATAAT GTCAAAATAG 60
CTATTTTCTC AATAAAAATC TCAATCTCC TGAAA 95

SEQ ID NO:7496

SEQUENCE LENGTH:143

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08738

SEQUENCE DESCRIPTION:

GATCCTCCCT CCTTGGCCCC CCAGTGTGG GATTACAGGC ATGAGCCACC GTGCTTGGCC 60
TGGAATNAAT TTTTATTGA AGTTATTCTG TATCTCCAGT GTTGGCATAT GCTAGACATT 120
CAATGAATAT TTGTTGATTT AAA 143

SEQ ID NO:7497

SEQUENCE LENGTH:342

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08739

SEQUENCE DESCRIPTION:

GATCAACAAA ATGTGGTATA TTCATACAGT GGAATATTAT TCAGCCTCTA AAAGGAAGAT 60
ACGCTGACAT GTGCTGCAAC ATGGATGANT CTTGAGGACA TGATGCTAAG TGAAGTAAGC 120
CAGTCACTGG AAGACAAATA CTCTATGCTT CCNTTATGT GAAGTATCTA GAGCAGTCAA 180
ATGCATAGAA ACAAGTAGAA TGGTAGTTGC CAAGGACTGG GGGAGGAGGA AATGAGGAGT 240
TGTTTAATGG GTATAGTGTT TCATTTTTCG NAGATAAAAA GTCCTGTGGN TTGGTTGCAC 300
AGTTAAGTTA TGTGANTGCT GTATGCCCAA CTCANCTGTT NN 342

EP 0 679 716 A1

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SEQ ID NO:7498
SEQUENCE LENGTH:236
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08740
SEQUENCE DESCRIPTION:
GATCCAGTCG GCCTGGAGAT TCTACGCCAC CAACCTCTCG CGCACAGACC TGCACTCCAC 60
GTGGCAGTAC TACGAGCGAA CGGTCACCGT GCCCATGTAC AGGTACCGCC GCCGGGCACC 120
TGCCACCAAG CAACTGTTTC ATTTTATTATT TTCCATTTGT TCTTAAACCC CACTTTTTGT 180
TGTTCAATTAT TTTGATTGAT TTTNNNTTCT TTAAATGTA TTTTTCACAA AGGAAA 236

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SEQ ID NO:7499
SEQUENCE LENGTH:160
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08741
SEQUENCE DESCRIPTION:
GATCCTAACC CTTTAGTATG CTGGAATTCT ACTCTTCACT TACTGCATTG ACTGTTGTTG 60
ATTAGTTATN ATTGCAAAGC ACTGTCACCG GCCTCAGGGA GTTTATGTGT AATAGAATTA 120
AAAATAATAG CTGTGTATAA CACTTAGCTC AAGCCACAAA 160

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SEQ ID NO:7500
SEQUENCE LENGTH:210
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08742
SEQUENCE DESCRIPTION:
GATCTTATTG TGGAGTTAAA CAAAGCACCC TTAAAGTAGC AAGCGAGCAA AATAAAAGCA 60
AGGAAAATCA GTTGGAACAT AACTTATGTA TTTTITAGGA TTAAGAGTAA TTAACTGAG 120
TCAATTCAAT ATGTTTTTAA AGGTGTGATT CTAAATNCT TTATGATTGT TTAGATTTCT 180
ATATCTTGGT AACATGAATA CACTTAAA 210

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SEQ ID NO:7501
SEQUENCE LENGTH:351
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08743
SEQUENCE DESCRIPTION:
GATCAGCATT TGTATTTNAG TCCATTAATG GGAAAAATAG CTGATATTGT NCATAGGAAC 60
TTGGAAACCA CACAGGACTT AAGTCCCTTG TCTGTNTTGA TGGTCAACAT ATCTCCTTTA 120
ATATCACGAC ATTTTCAACA ACANCTGGTG AACAAAACAG ANCTNCTTTT TGACACCATA 180
GATTCTCCTG AGGTCAACGT TGCAAAANGC ATAGCAAAGT TNCTCCGAAA TGTTAGATAT 240
CGTTATCAAC CACTATTAGA CAGATGTAAT ACCGTATTTT TTCAGTAATG TGGNCCACCT 300
TGATTTTGGN TTCCATCAGT ANAGNGACTT AGTGTATACA NNNACNAAAN N 351

SEQ ID NO:7502

SEQUENCE LENGTH:195

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08744

SEQUENCE DESCRIPTION:

GATCGCACCA CTNCACTCCA GCCTAGGTGA CAGAGGGGAT GACAGAGGGT GCTCTGCGNN 60
CAGGGAAGGC GGGGTGGTTG GGGGGTAGAG AATGTTGCNC TNATAACCAC AGCTGTGGTT 120
ATTTTNTGGC AGGTACAGGT NAGGTTCCCT CTCCTTTCCA CCAGATGGCA CTGGAGAAAA 180
GGGATTNNAN TTAAN 195

SEQ ID NO:7503

SEQUENCE LENGTH:257

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08745

SEQUENCE DESCRIPTION:

GATCAGAGCA GAAATAAGTG AAATAGAGAC TAAAAAATAG AAAACATCAA TGAAACAAAG 60
AATTATTTTT TAAAAAGATA AACTAAATTG ACAAACCTTT AGCTAGACAA AGAAAAAAGA 120
GATAAGACAC AGATAANTAA NTTCAGAGAT TAANAAGAAG CCATTATACC TGATACCACA 180
GAANTACAGA GANTCATAAG AGACANTTAT GTTGTGCACA TGTACCCTAG ANCTTAAAGT 240
ATAATANNCN NTANTTN 257

SEQ ID NO:7504

SEQUENCE LENGTH:328

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08746

SEQUENCE DESCRIPTION:

GATCTTACAG AGAAAGACTA TGAATACTT TTCAAATCTA TTAATGGAAT CCTTNCCT 60
GGAGGAAGTG TTGACCTCAN ACGCTCAGAT TATGCTAAAG TGGCCAAAT ATTNATAAC 120
TTGTCCATAC AGAGTTTTGA TGATGGAGAC TATTNCCTG TGTGGGGCAC ATGCCTTGGA 180
TTTGAAGAGC TTTCACTGCT GATTAGTGA GAGTGCTTAT NAATTGCCAC AGATACTGTT 240
GACGTGGCAA TGCCGCTGAN CITCACTGGA GGTCAATTGC ACAGCAGAAT GTNCCAGANT 300
TTCCCTACTG AGTTGTTGCT GTCATTAN 328

SEQ ID NO:7505

SEQUENCE LENGTH:102

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08747

SEQUENCE DESCRIPTION:

GATCAATTTG AAGTCCTGT TTGGGAATAA GGCACCTATC AGCATGAAGA ATTTTTTCTC 60
ATTCTGTGCC ATTTAAAAA TAGAATACAT TTGTATATTA AA 102

SEQ ID NO:7506

SEQUENCE LENGTH:343

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08748

SEQUENCE DESCRIPTION:

GATCACCTTC GNACGACTAC ATCGCCTGCT GCGTCAAAC T GAGGGCTCTT ACAGNCAGTT 60
 TTCGAAGNCG GGGTACTGCT CAGCAAGGTG TTGTGAATTT CCCATATGAT GATTTTCATT 120
 AATGTGTTCAT GAGTGTTTAA ATCAAGAGGA AGCTGCATGA ATGTAATCAA CATTCCAAC 180
 GGAGCTCTCC TTTGCTTGTC CTCCTTGCCT TCGGTAATAT GTATAAACTT ACATCACGAC 240
 TTTCTCTTAA CAGCTGTTGT AAAGTTTATT ACTTTATGTA CAACTGAAGT TTTGTTTTAG 300
 TTTTGATAAT AAAATTCTTT GGAACCTTAA AAAAAAAAAA NGN 343

SEQ ID NO:7507

SEQUENCE LENGTH:342

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08749

SEQUENCE DESCRIPTION:

GATCCAAGTG CTTNTTTGGT ACCAAGAGAC CCGGNTCTNT GCCAGCATAA ACCACTCACA 60
 CCCCAGGGGG ATGAGCTCTC TNAGCCCAGG ATTCTGGCAA GGGAGGTGAA GAAAGTGGAT 120
 GCGCAGAGTT CGGCTGGGGA AGAGGACGTG CTCCTGAGCA AAAGCCCATC CTCACTCAGC 180
 GCTAACATCA TCAGCAGCCC GAAAGGTTCT CCTTCTTCAT CAAGAAAAAG TGGANCCNGC 240
 TGTCCTCCCA GCAAAAACAG CAGCCCTNAA TAGCAGNCCN CGGTCTTTGG GGAGGNGCAA 300
 AGGNAGGCTC CGGTTGCCCC AGATTGGCAG CANNATTATN AN 342

SEQ ID NO:7508

SEQUENCE LENGTH:369

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08750

SEQUENCE DESCRIPTION:

GATCAAGCAA TTAATCTTCT GTAGAACCAA AATTNACTGT AATAACTTGG AGCAGTACTT 60
 TATACAACAA GGAGGAGGAC CTGATAAAAA AGGACACCAG TNCTCATGTG TTTGTNTTCA 120
 TGGTGACAGA AAGCCTCATG AGAGAAAGCA AACTTTGGAA AGATTTAAGA NAGGNGATGT 180
 AAGATTCTNG ATTTGCACAG ATGTAGCTGC TAGAGGNATT GATATCCACG GTGTTCCCTTA 240
 TGTNATANNT GTCACCTGNC CCGATGAAAA GCAAACTNC GTACATCGAA TTGGCAGAGT 300
 AGGGTGGAGC TGANGGATGG GTCTGGCAAT TNCCCTNGTG GCACCGAAAN NGGACAGGGT 360
 TTNGTACCN 369

SEQ ID NO:7509

SEQUENCE LENGTH:335

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08751

SEQUENCE DESCRIPTION:

GATCTGGCCA GAGGGGTGTT GGAGCCCAGC CCACCCACAT ACCAGTCAAG CTCTTAGGGG 60
 AGCAGAAGAA AAGCAGGAAG AATTAAATG TTTAATTTT TTTTNAAT GACTTTNCTA 120
 GTTATTAATA GTTGCTTGTT TCAGCAGTGN TATTGTATAA NGACCATCTT GTAAGNTACT 180
 CCTGACATCT TGCITTAGCA CATGTACAGT ACAGTTNCTN TGATAATGTG TTTGCCCTAA 240
 CTNCCCTGGC TTCNCCTTCA GCCCATCCNC TCTCCTCTAG NGCAGTTGGG TTGGAGGCTC 300
 NTTGAGGCAA GCNGCANCAT TGGNGGGGGT GCAGN 335

SEQ ID NO:7510

SEQUENCE LENGTH:270

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08752

SEQUENCE DESCRIPTION:

GATCCGACAC AGAGAACAGG AGGAACTNAG GCGACAGCAA GAGGGCTTTA AGCCAAACTA 60
 CATGGAAAAT AGAGAACAGG AAATNAGAAT GGGTGATATG GGTCCCCGTG GAGCAATAAA 120
 CATGGGAGAT GCGTTTAGCC CAGNCCCTGC TGGTAACCAA GGTCTCTCTC CAATGATGGG 180
 TATGANTATG AACANCAGNG CAACTNTACC TGGTCCACCA ATGGGTACTG GTNCTGCCAT 240
 GGGACCAGNA AGGNGCCGCA ANTNTNGGGN 270

SEQ ID NO:7511

SEQUENCE LENGTH:85

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08753

SEQUENCE DESCRIPTION:

GATCCCAGGA GAATCAGAAA CTCCAACATT TTGGAATCTT CAAGGGCACA TACTGAGAAA 60
 AAAAAATAAA TTGTTTATGA GCAAA 85

SEQ ID NO:7512

SEQUENCE LENGTH:66

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08754

SEQUENCE DESCRIPTION:

GATCAAACTA TACGCTGTCT ACAAGAGACA CTTTAGATTT AAAGACACAA ATAGGTTGAA 60
 AGTAAA 66

SEQ ID NO:7513

SEQUENCE LENGTH:390

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08755

SEQUENCE DESCRIPTION:

GATCTTTTGT ACTGAGGTTT TTAACACTTT ACTTGGGTTT ACCAAGCCTC AACTGGACAG 60

EP 0 679 716 A1

5 ACCATAAACA GTCCACAGGC ACCGTTCTTG CCAGGCCCCA ACCCAGAGG AGTCTNTCCG 120
 CAGAGCCTTC TTGGTGTTGC CCTAACTTGC CAGTGGCCTT TGCTCAGAGC CTCCTCCTGT 180
 GACATGTGAA CAATGAAGAG GCCTGCGCCT CCTGCCTTGC CGCCTGCAA GCAAAGAAAC 240
 TGCCTTTNAT TTTTAACTT TAAAAAGTAG CCAGATTAGT ANCANGACTG GCTGGCTTGA 300
 TGAGCAAAGC CTTTGCTCTC ACGGNAGGAG GGTAGGGTTT NGGNTGTACA ATGGTAACTT 360
 GCNNTGGGNA CCTTAAAAAG GGAGGTGNAN 390

10 SEQ ID NO:7514
 SEQUENCE LENGTH:212
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08756
 15 SEQUENCE DESCRIPTION:
 GATCACCTGA GGTCGAGAGT TTGAGACCAA CCTGACCAAC ATGGAGAAAC CCCGTCACTA 60
 CTAAAAATAC AAAATTAGCT GGGCGTGGTG GCACATGCCT GTAATCCAC CTACTTGTGA 120
 GGCTGAGGCA GGAGAATCAC TCGAACTCAG GAGGCGGAGG TTGNAGTGAG CCACTGCACT 180
 20 GCCTGGGCAA CAGAGCGAGA CTCCATTCA AA 212

SEQ ID NO:7515
 SEQUENCE LENGTH:136
 SEQUENCE TYPE:nucleic acid
 25 TOPOLOGY:linear
 CLONE:HUMGS08757
 SEQUENCE DESCRIPTION:
 GATCCACCCG NCTTGNCCCC CCCAAAGTGC TGGGATTACA GGTGTGAGCC ACTGTGCCCC 60
 30 GCCGAGTTAA ATATATTCT AACATAGTAA AAAAAAAG GTANGTTNCA TTAAATAAAT 120
 CCNGGAAGTN CAAATN 136

SEQ ID NO:7516
 SEQUENCE LENGTH:160
 35 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08758
 SEQUENCE DESCRIPTION:
 40 GATCATGAAG ATGGACAGAA CAGACACTGT GGCAATAAGA TACCAAATTG TAAACAGGAC 60
 CTAAGGCCAT GCCAAGCAAG GGTAAAGTCA TGTAAGAGCC GCCTTCATGC CCCGCCTCCC 120
 CTGCCCCCAA CTTATGAATA AACTATGTTT TAACTGCAA 160

SEQ ID NO:7517
 45 SEQUENCE LENGTH:355
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08759
 SEQUENCE DESCRIPTION:
 50 GATCTGGAAC TTGAAGATGC CATTATACA GCCATCTTAA CCCTAAAGGA AAGCTTTGAA 60

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EP 0 679 716 A1

GGGCAATGA CAGAGGATAA CATAGAAGTT GGAATCTGCA ATGAAGCTGG ATTTAGGAGG 120
 CTTACTCCAA CTGAAGTTAA GGATTACTTG GCTGCCATAG CATAACAATG AAGTGACTGA 180
 AAAATCCAGA ATTTCAGATA ATCTATCTAC TTAANCATGT TTAAAGTATG TTTTGTTTTG 240
 5 CAGACTTTTT GCATACTTAT TTCTACATGG TTTAAATCGA CTGTTTTTCA AAATGNCACT 300
 TATANNTCCN AATAANCTGT TAANCCNGGN ANANGGNNTT GTNTTNGTGN GGNTN 355

SEQ ID NO:7518
 SEQUENCE LENGTH:77
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08760
 SEQUENCE DESCRIPTION:
 15 GATCTTCATT TTCCTGTTCT GTATTCTGCT ATGAAGTTCT CTGTTTTTCA AATAAAATTT 60
 ATGTGTAAA TAACAAA 77

SEQ ID NO:7519
 SEQUENCE LENGTH:20
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08761
 SEQUENCE DESCRIPTION:
 25 GATCATCTTG GAACTGCAAA 20

SEQ ID NO:7520
 SEQUENCE LENGTH:247
 SEQUENCE TYPE:nucleic acid
 30 TOPOLOGY:linear
 CLONE:HUMGS08762
 SEQUENCE DESCRIPTION:
 GATCGGAGGC TGCAGTGTGA CAGTGCCAGC CAATGTGCAG AGGTGGATGA GGTCTTGTA 60
 35 AAACCTGGCT CCTTTTAAAC CGGCCCTCAA GCTCCTTAAG TNAATTGCCG TAACTAATTT 120
 TAAAGGGTTT AGATTTTAAAG AATGGTGCTC TTTCATGCCT ATNATCAGTA AGGGGACTTG 180
 TATTAGAGTG AGAGTCTTTT TATTTAGGCC AGTTGTCAAG TGTCAATAAA AGCATCATGT 240
 AATTAAA 247

SEQ ID NO:7521
 SEQUENCE LENGTH:164
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 40 CLONE:HUMGS08763
 SEQUENCE DESCRIPTION:
 GATCTTAATA TTGACTAGTT TCACATCCAG GTTTCTAAGA AATGATAAGA TACTTCACTT 60
 TTCCAGAGTG AAATGTAGGA GGGAGCACAT TCTAAGTACA GCTAAAAATT TAGCTCACTG 120
 50 TAACACAGTT TCACTCTCTG AATAAATAAA GCAAAAAACA CAAA 164

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EP 0 679 716 A1

SEQ ID NO:7522

SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08764

SEQUENCE DESCRIPTION:

GATCTGCCCTG CGAAGCTCCC CACAACACTG TCCCTGGAGA CCACCCATGT GGCAGACACT 60
CAAAATCTGT TGTATTTGCT AAATAAATTA ATATGGACAA GTTACCCAAA 110

SEQ ID NO:7523

SEQUENCE LENGTH:181

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08765

SEQUENCE DESCRIPTION:

GATCAAATTG TTCATGCCCTT GGGATTTTTT CTTCTATAAT TTGGATATCT TTTCCATTG 60
CTCATTTCTT TATTGAAGTT GCTTTCTTAA AAATTGATTT AACTTTGTCC TTGGTATCTT 120
TAGTTCAGTC CAGAAATNAT TAATTGTGAT GTGATTAAAT TCATTTTTTG GTCTGGTGAA 180
A 181

SEQ ID NO:7524

SEQUENCE LENGTH:175

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08766

SEQUENCE DESCRIPTION:

GATCTGAATG GTTTAATAA GGGGCTTTCC CCCTNCTCG CTCCTGCACTT CTCCTTGCTG 60
CCATGCGANG AAGGATGTGT TTGCTTCCCC TTCTGCCATG ATTGTAAGTT TCCTGAGGCC 120
TCCCCAGCCA TGTTAACTT TGAGTTAATT AAACCTCTTT CCTTTATAAA TAAA 175

SEQ ID NO:7525

SEQUENCE LENGTH:362

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08768

SEQUENCE DESCRIPTION:

GATCTGAAAA ATNCTTTTTG TCCTCATTGT TGCAGTTTCA ATTGTAATCC ATTGCATCTN 60
ATTTTCTTGG GTGTTGGCAC TGTAATATAT TAAAAAATAA TTCAGTGCTC AGTATTGTAA 120
CAACTGTCCC TACTGAATAT TCCTTTTCAT AAATACTTGC TACCACAGGG AAATTAAGTT 180
TTCATATAGA ATAAGTAGTT TCAGTAGTAA CCATTGAAGA NTTAANCNTG TGGTTCANGG 240
CAGATATTTT TCTGGAAAGT TTTCTCTATT GTANAATTTG TNGTATATGG CTATGCTGAC 300
TATCNTGGAT NCTGTCATGG GACATGCATT ACCACANNTN CNNGCCCCTT TCTGTGGTNN 360
NN 362

SEQ ID NO:7526

SEQUENCE LENGTH:306

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08769

SEQUENCE DESCRIPTION:

GATCCCATGG ACATTG GGG AAAGGGCTCC TTGGGCTGCT GGTGAAC TTC TGTGGCCACC 60
 ACCTCCTGCT CCTGACCTCC CTGGGAGGGT GCTATCAGTT CTGTCCTGGC CCTTTCAGTT 120
 TTATAAGTTG GTTCCAGCC CCCAGTGTCC TGA CTCTGT NTGCCACATG AGGAGGGAGG 180
 CCCTGCCTGT GTGGGAGGGT GGTACTGTG GGTGGAATAG TGGAGGCCTT CAACTGATTA 240
 GACAAGGCCC GNCCACATCT TGGAGGGCAT CTGCCTTACT GATTAAAATG TCAATGTGAA 300
 TCTAAA 306

SEQ ID NO:7527

SEQUENCE LENGTH:76

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08770

SEQUENCE DESCRIPTION:

GATCTTTTAG TTATCAA ACT GTTAAATACA TTTTAACTTT CAAATTTTAT GAATAAAGGA 60
 GATGAATCCT GTTAAA 76

SEQ ID NO:7528

SEQUENCE LENGTH:74

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08771

SEQUENCE DESCRIPTION:

GATCTGAAGT CTGATTGTGT GTACGTTGCG GCGTGCGTTA GTCCCTGCCA CGATTAAAGA 60
 CTTAGACCGN CAAA 74

SEQ ID NO:7529

SEQUENCE LENGTH:102

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08772

SEQUENCE DESCRIPTION:

GATCGGCCA CTGCACTCCA GTCTGGTGAT AGAGCAAGAC TCCATCTCAA AAATAAAAAA 60
 TAAAAATGTTG GNTGTGGGGA GCCAGGCCTC TCACAACTGA AA 102

SEQ ID NO:7530

SEQUENCE LENGTH:312

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08773

SEQUENCE DESCRIPTION:

GATCGAATCC AGGCCCTGGG GCTTCANTTG CCAGACCCAC CATTAGAGAG TGAAGATGAA 60
 GATGAGGAGG GAGCTACAGC GTTGAACAAC CACAGCTCTA TTCCCATGGA CCCAGAACAT 120
 GTAGAGCTGG TGAAGAGGAC AATGGCTGGN GTAAGCCTGC CTGCGCCAGG GGTTCCTGCC 180
 TGGGCTCGGG AGATATCGGA TGCCCACTGG GAAGATGTGG TACAGAAAGC CCTCCAAGCC 240
 CGGCAGGCAT CCCCTGCCTG GGAGTGACCA CAGTGAGAGC TGCCTTATAT TCCTACATTC 300
 CAGGCCAGAA CN 312

SEQ ID NO:7531
 SEQUENCE LENGTH:186
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08774

SEQUENCE DESCRIPTION:
 GATCTGGACT GAGGACCCTG CTAATCCCCA AGCCAGAGCC CATCAGCCAG GCCTGCTGTG 60
 AGCCACCTGC CTGTGGAGTG CTGAGCTCAA CCAAAGGCTG GCAAGCTCTG GGCCTCATTT 120
 AAGGGATTCT GATGAGCCGA TGGGCCCTGG AGGCAGCCCA TTAAAGCATC TGGCTCGTTT 180
 TTGAAA 186

SEQ ID NO:7532
 SEQUENCE LENGTH:390
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08775

SEQUENCE DESCRIPTION:
 GATCACATTG CCATTCTCC CCTCCAGAGG TAACAATTAT CCACAATTG ATGTTTATCA 60
 TTCCTGTGTT GTTGTACTTT CACTGTGTAT AACCTAANNC NTCTACTNTT TAGTACTGTT 120
 TTATATATTT TNAAGCCTCA TACTTGCTCA TTCTACAGCT TTTTCACTC ATTATTGTAT 180
 AATTATATCT GANGCNCTCG TTCATTAATT TTAGTCCNGT GTAGCAGAAT TCAATTACGG 240
 GAACTACCAT AATTTATCTG TNCTCCAGTT GAAGGCATGA NGTTGTTGCC AGTTTCTGTA 300
 TTATAACCCAC TGTAGTGGGA CCATTCTTCT GCNTTGGGCT CACTGCGGGT TACCNAAGNC 360
 GGTNCCNCAG GATAANCCCC ATTTGGGNTN 390

SEQ ID NO:7533
 SEQUENCE LENGTH:378
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08777

SEQUENCE DESCRIPTION:
 GATCTAGCAT GTGGATTTTA AAAGATTGCT CCTCATTAAAC AAGAATAACA TTAAAGGAG 60
 ATTGTTTCAA AATATTTTNN CAAATTGAGA TAAGGACAGA AAGATTGAGA AACATTGTAT 120
 ATTTNNCAAA ANCAAGATGT TTGTAGCTGT TTCAGAGAGA GTACGGTATA TTTTGGTAA 180
 TTTTNTCCAC TAGCAAATCT NGATTNNGTT TGATAGTGTG TGGAATTNCC TTTTGAGGGA 240
 TAAGACCATG GGANANTTGT GGTAAAGACT GTTGTNCCC TTCATGANTN CTTCTGAAAT 300
 TGCCNTCAGT TTTNCTAATC TCCTGTGAAA TGCCTAGGTT TTGCGCCTGT CCCACTTTTT 360
 NTTTGTGGGN CTTATNTN 378

SEQ ID NO:7534

SEQUENCE LENGTH:175

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08778

SEQUENCE DESCRIPTION:

GATCATTGA GTGAAAAA GGTACCTGTN CAAGCAAGCT CCTGGACACC ACAAGAAGGA 60
GGAATTATTT TAAAAGCTGT ACTCTAAAT TGTNAGTATC TTAAAAATCA GTTGTGAACA 120
ATGAAGGATT TGAAAGAGCA TTGACTTTGC CACTTAAAAG TATTTCTAAA ATAAA 175

SEQ ID NO:7535

SEQUENCE LENGTH:159

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08779

SEQUENCE DESCRIPTION:

GATCAAGGTG AATTTTTGT ATCACTGCCT GTGGAAATCT ATCCTCATCA GTNATTGCAT 60
TTTCCCTGC CTATACCTGT GCTCCTTTT MTNACTGTGT TTTCAGTCAC TTCCTTCTG 120
TNAAGGTTG CTTAGCTTTT NTTTTNAANA TTTGTTGTN 159

SEQ ID NO:7536

SEQUENCE LENGTH:304

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08780

SEQUENCE DESCRIPTION:

GATCCAAACC CACAGTNTTG TCCTGGAGGC AGCAGGGGTG AAGGTGGAGG GTCCAGGGCC 60
ATGAGGAGCC CCCTTGCCAT CAGAGCCTGG CCTAACCACC CTCCTCTCTA CTTACACACA 120
CATGCATTTT ATAATAGCTC TGACCCAACC TGGCCACTCT GCAGAGACTG GGACAGACAG 180
GTGCAGGCAA TGGGCCCTCC CACACCCAGT CACCTACAAG GAATTTTCAA ATCCACTTTT 240
AAAACAGAAA CCGGTAAATG CGCCGTATTG TATATTTTAT TTAAATAAAA AAAATTCCAG 300
CAAA 304

SEQ ID NO:7537

SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08781

SEQUENCE DESCRIPTION:

GATCCACCCT TCCTGGGCCT CCCAAAGTGC CGGGTATTAC AGTCATAAGC CACCACACCC 60
GGCCTAACAT GCATTTTTTT AATTCAATAA ATATTATTG AGCACCTAAA 110

SEQ ID NO:7538

SEQUENCE LENGTH:160

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08782

SEQUENCE DESCRIPTION:

GATCCCCTCC CCTCCTGCCT TTCTGTATTT TTTAAAAATT ATTTTGTTTT TAATTCACAA 60
 ATATTGTATA TGTTATATGG GGTAGAATGT GATGTTGATA TATGTTTAAA ATGTGGAATG 120
 ATTAAATCAG GCTAATTTAC TTTTTTNNCC NAAAAGGAAA 160

SEQ ID NO:7539

SEQUENCE LENGTH:219

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08783

SEQUENCE DESCRIPTION:

GATCCACCCA GTTGCTCAAG TAAAAACCT GAGAGACACC TTACGCCATC TGGGAACCTG 60
 TTAGAAATTC TTAACCTCA TTCTATACCT GCTAAATCAG AAAATGTAGG GATTAAGTCT 120
 GATTATTTGT GTTTTCGCAA GTCATCCNGG TGATTCTGAT GCATGGAAAG TTTGAGAACC 180
 ACTGGTTTAG GTTATTAATA GTTCTGTCT CCCTATAAA 219

SEQ ID NO:7540

SEQUENCE LENGTH:355

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08785

SEQUENCE DESCRIPTION:

GATCAGAGAA GACTTTAAAA CTCTTGACTT AATTGAGTAA ACTCTTCATG CCATATACAT 60
 CATTTCCTT ATGTAAAGG TAAATATGC TTTGTNAACT CAGATGCTG TAGCCAGGAA 120
 GCCAGGGTGT GTAAATCCAA AATCTATGCA GGAAATGCCG AGAATAGAAA ATATGTCAC 180
 TGAAATCCTA AGTAGTTTGT ANTTCTTTG ACTTGAATCT TACTCATCAG TAAGAGAACT 240
 CTTGGTGTCT GTCAGGTTTT ATGTGGTCTG TAAAGTNAGG GGTCTGTTT TNGTTTCCTT 300
 ATTTTGAGGT AAGAGTACTG CTNGGTGGTC GGGGGGGTTA TATNNTNCCC TTTTN 355

SEQ ID NO:7541

SEQUENCE LENGTH:314

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08786

SEQUENCE DESCRIPTION:

GATCAGGTTT CCCCTCAGAA CCTGTGTTGC CTGCCTGCTG TATCTTCCAT GTCTTCTTA 60
 TTCAACTTCT AGAAGTAGAG AAGGTTGGGA CTTGAGTGTA GAAGAAACAT CAGAATAATT 120
 TCAAAAAACA GGAAGTTGGA ACTATAATTT TAAATTATT GTATTGTTTT TACGGATTTT 180
 ACATGTATAG TAAATATTA ANNTTCATCT CTTGCAGTTC ANTAAGCT ATAGTATAGA 240
 TTTTATTTT AAGAAGTTA TATAATNCTT TTGTGAGCCA ATATGTAATA CTNGAATATT 300
 TATTTANCT ANTN 314

SEQ ID NO:7542

SEQUENCE LENGTH:298

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08787

SEQUENCE DESCRIPTION:

GATCACAGAG AAAGTGATTA CAGACATTGC TGGGGTAATA CCAGCTGAGA AAATTGATGG 60

AGTATTTNCT GCCTGTNAGA GTGGCTCTTT TGACAACTA GANGCTGTGG TCAAGGATTT 120

AATAGATGAG GGTGATGCAG CAACTCAGCT CGTCAATCAN CTCCATGATG TGGTTGTAGA 180

AAATANCTTA TCTGATAANC AGAAGTCTNT TATCNCAGAA AACCTTGCCG AGGTTGCCAA 240

ANGCCTGGCA GATGGTNTCG TTGTCCATTT GCANCTCATC NGCCTTTTNG CCCTGNGN 298

SEQ ID NO:7543

SEQUENCE LENGTH:32

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08788

SEQUENCE DESCRIPTION:

GATCAAAATT AAAGACAGTT AAAAAGGTGA AA

32

SEQ ID NO:7544

SEQUENCE LENGTH:249

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08789

SEQUENCE DESCRIPTION:

GATCATCATT TATTGCATCT CATAACTAAA TTTNNTAAAG TTTGGATTGG GACTTTTNAG 60

GTCCTTTTGG GAGGGCAAAG GAAGTGCCAG CTTCTNTGGG GAACTTGTTT TTAAATCCAA 120

AGACTTGAAC CACATTCCCT GCACATGANC ATGTTTGCTT TTATCCCTTC TCTCATTGTN 180

TCCTACCCAT CTTAGTACCA TTGTGGTTAT AANCATCTGC ATTTNTNAGA AGCATTTNAC 240

CCATTTNNN 249

SEQ ID NO:7545

SEQUENCE LENGTH:257

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08790

SEQUENCE DESCRIPTION:

GATCATATGG AATGTACTAT TTTGTAATGT CTTTTTTCAT TTACAATGT ATTATCAACC 60

TTTTCCCTCT CAAAAATACA TTGTGAATGA CTGCATAGTA TTCACTTTAT GANTATTTAA 120

TTCATTTTAC AGTCTTCTAT TGTNGGACCA CTNACATTGT ACCAAATGTT TTCCTTTGGT 180

TTATTCTTTA ATGTATTAGT ATTTTACTGC TGGTCACTCA TGGAAATCCTG CAGCTTTANT 240

NAAAAGCAAG GATGAAA 257

SEQ ID NO:7546

SEQUENCE LENGTH:158

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08791

SEQUENCE DESCRIPTION:

GATCGTGCCA CTGCACTCCA GCCTGGGCGT TGCAGTGAGA CTCCGTCTCA ACAACAACAA 60
AAAGGTGGAT ACATGTCAAA AGGGCACAGG AGCCACTTGA AGGATTATAC TTGCTAAATC 120
TGTGTCAATT ATCAAAATAA ACTGGTAGCA ACGTTAAA 158

SEQ ID NO:7547

SEQUENCE LENGTH:128

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08792

SEQUENCE DESCRIPTION:

GATCAAAGGG ATGAGCAACA GGGACTTCTG CCACAGTGAC AATGGAATTG TGTGTGCCT 60
TACTTCAGAG GTGGTCTCTN CTTTCTTGTA ATAAAAGCAA TATTATGCG GAAAGCAAGC 120
AGCTCAAA 128

SEQ ID NO:7548

SEQUENCE LENGTH:205

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08793

SEQUENCE DESCRIPTION:

GATCACATCA AGCTAAAAA CTTCTACACA GCGAAGGAAA CAAAGTGAAC AGAATAACAT 60
GGGAATGTTT TCTGTAATTT AGTAGTAAGT GGCAATAGTT TACAAACACA TTTTGTGTAT 120
ACTGCTGTCA TTGCACTGNT TACCTTCTGT TGTAGTGACT TTGTTCTATT AGTCCACTCA 180
ATTAAAAATAT TTGGTTTTGT TAAA 205

SEQ ID NO:7549

SEQUENCE LENGTH:372

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08794

SEQUENCE DESCRIPTION:

GATCTTATTT TTAATGTAGT GAGGATATTA TTAAACTTT TATTTAACT GGAAATGTCC 60
TGAAACACAT ATTTAAATA TTGGGATACA GTGAAAGAAA AATTCAAATT TTAATAACAT 120
AAAGATTTC TAACTTTATG TTATTGANCA CTTACTCACT AGAAGTGAGT TCTTTAGAAA 180
ANTACAGTGA AGGACTCAGT TCAGTCTTGT TTTTATCAGA GTGATAATCA TCCTGTTTCA 240
CATCCCAATA CTATTTTGAN ATTCTAAACA ATGGAACCCA AANTNCCAAT AANTATAAGG 300
TTATGCCTTC AATATATTCC TATACANTTC TGTAACCATG GTTTANNATA CACAAGCTTA 360
NNATACCATG GN 372

SEQ ID NO:7550

EP 0 679 716 A1

SEQUENCE LENGTH:97

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08795

SEQUENCE DESCRIPTION:

GATCCGGTNG TCTAAATGCA TTCATATTTT TATGATTGTT TTGTAAATAT CTTGTATAT 60
TTTCTGCAA TAAATAAATA TAAAAAATT AGAGAAA 97

SEQ ID NO:7551

SEQUENCE LENGTH:244

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08796

SEQUENCE DESCRIPTION:

GATCTGGGTT TGGCATTAGA AGTATTTTCAT AATTTTGGTT TTTNATTTAG GTTTCCTCAA 60
CATCTGTAAA GTGATTGATT AAATTAGAGG AGGCGGTAG AATAAATCCC AATCCCATTG 120
CAACTGGCAG AGCATTATAA NTNTTTATAA ATNCAGTTAC AACAAAGGAG AGGNTCTACA 180
TTCTTCAAAA AANAAANTTG GCTTACCTGT TTCANCTTN GGTGGCATT NNCTGCCATT 240
NGTN 244

SEQ ID NO:7552

SEQUENCE LENGTH:364

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08797

SEQUENCE DESCRIPTION:

GATCTGGGAA TTACCCATCA TCATTAAGCA ATGAAACAGA CCGGCTGGTG CACTGGTGCC 60
ACGGCGCCCC GGGGGTCATC CACATGCTCA TGCAGGCGTA CAAGGTCTTT AAGGAGGAGA 120
AGTACTTGAA AGAGGCCATG GAGTGTAGCG ATGTGATTG GCAGCGAGGT TTGCTGCGGA 180
AGGGCTACGG GATATGCCAT GGGACTGCTG GCAACGGCTA TTCCTTCCTG TCCCTTTACC 240
GTCTCACGCA GGATAAGAAG TACCTCTACC GAGCTTGCAA GTTTCAGAG TGGTGTCTAG 300
ATTACGGAGC ACACGGGTGC CGCATTCCTG ACAGACCCTA TTCGCTCTTT GAAGGCATGG 360
CNAN 364

SEQ ID NO:7553

SEQUENCE LENGTH:129

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08798

SEQUENCE DESCRIPTION:

GATCTGTGTG TCTCTGTTTT GAGTTTTTCC TGTTTATTTT GAAAAGTACT GTTGGTCAAG 60
ATAATTGGTC AATAATCCAT GTTGGTTTAA AAAAAAGCA TTTTAACATT AAAAATATTA 120
CAGTATAAA 129

SEQ ID NO:7554

EP 0 679 716 A1

SEQUENCE LENGTH:316

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08799

SEQUENCE DESCRIPTION:

GATCACATTT TNATTTTTTN TTCTCCACAT CAGGATAGTT TACTGAAGCA CAATCTCTTA 60
TACTAGTGGG ACAAAGGGA GAAAAGGAA GCAAGATAAA TGGGTATGTA GGATGAAGGG 120
TTATTTAAAA TGGAAGTAAA GATAGAAGGA GGACTGTAGG AAGANATGGA ATNNTTTANN 180
TNTGAGGAAA GATATCTNTG GTAGACATGT CCTTCCATGA CTAATTTCTA ATTGTNACTC 240
ANCAACACATT GAGGTATGGN CCCTCCTCAG TGACTTTNAC TAGCTCAGAN ACGTACTCCC 300
CCACCANCCC CACCTN 316

SEQ ID NO:7555

SEQUENCE LENGTH:365

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08800

SEQUENCE DESCRIPTION:

GATCAGTGGA ATTCTTGGTT TGATGATAAC CTTATTAATT GAAATNTTTT ACTGATGTGG 60
CTTTAAAAGA GGTATTATNTT GTATATGTTT AGAACTCTCT GATTTTATG AATTATATGG 120
GAATGAGAAA CAGAAGAAGT GGTATTTNCT GGCGAGTTAA ATAGGCAAGG TACCCAGTGA 180
TAACACCAAC CAAACCACTC CTATCTGCAT GATTCTGAAC ATCTGGATGC CTGTGGTTTT 240
ACTGTGTATA TNTNATTTTT AATATATTAC CTTTGTGGAT NCATTTAAGG TCTACTCAA 300
NGTANCACTG TCAAAACCAC TAATATGTAT GTNAAANTTG TNCTGTNTAC TACATTACNG 360
TNCAN 365

SEQ ID NO:7556

SEQUENCE LENGTH:349

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08801

SEQUENCE DESCRIPTION:

GATCTTGAGT AACTGAACTT ATTTCCCTTA AGGTATAAAA ATTTAAATAA CCTGTATATA 60
TTTGNTAACA NCTTTATNAG GGAGGTAAGT NACTTACAAT ACATTGCACA TATTTAANGT 120
ATAGANTTCG ACGAGTTTTG ATGTATGTTT GCACCTGTGA AACCATCACA ACAAACAGTG 180
AGCATATCTN TCACCCCTAA NNGTTTCCTC CTGTCCCTTG GTAATCCCTC TTTTACCTGT 240
CCCACCTCTG TTCCTCAAGA NCCACTTACC TGCTTTNNTN ACTGTTAATG TACGTGAGTT 300
TACATTTTTG AGAGTTTTAT CCAAAGGNA TCATGANGTA TGTNCTCTN 349

SEQ ID NO:7557

SEQUENCE LENGTH:369

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08802

SEQUENCE DESCRIPTION:

5 GATCTNATCA ATCTAGCGGG AGAGACAGGA TAACCTGTCC GAAAGTATAG CGCCACTATN 60
 ACTCCGCCGG AAAAATTACT TTAAAAATCG CCAAAAATTA CTTGGAGCAA AGGGCAGTCG 120
 GCGGANTTCG CCAAGGCTGG CGCAGTCGGT TTTNACCTGT AGCAGAGAAC CNGGNCTGGA 180
 GAACAGCCTC ACTTCTTTNA TTGANTACTT ACATAATGCA TTGGAACATG ACATGAGATT 240
 AAGGTTTAAT AATGATAGAN TGAAGACCAC AATAAAGAG ACCTCTACTT AGCTCAGCAA 300
 TTCTTACCTT TCTTACCTAT TTGATGGAGG NTGTCTTTTG ANAGGTGTNC TGCAGGGNCC 360
 AAANTGTTN 369

10 SEQ ID NO:7558
 SEQUENCE LENGTH:290
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 15 CLONE:HUMGS08803
 SEQUENCE DESCRIPTION:
 GATCTTGAT GCCAACTTT AATCTGCTTT TATGTTTTCA GGCTGAAGGT GTGAAAATCC 60
 TAAGAGGATT TCATATTGAA TATGTGTACA CAATCTTAAC TATCGTGGTG GAAAACATAC 120
 20 TACTATAATT NATTATTATA TCTNCCAGAT AATGTTATTC ATTTAGAACA AATAAGGTAT 180
 ATTTNTNAGA ATCANCTTTG TAAGCACTAT AANNCTTTN ATAAGTTATA AGGTCTATGA 240
 TGTGTTTACT TTAANNCTTG CTGTTAANAG CAACACGTAT TANNTATGTN 290

25 SEQ ID NO:7559
 SEQUENCE LENGTH:55
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 30 CLONE:HUMGS08804
 SEQUENCE DESCRIPTION:
 GATCTGGATA AATTTTCATT TCTACTTAAT TAAAACTTCC TATGTAAAAT CAAA 55

35 SEQ ID NO:7560
 SEQUENCE LENGTH:109
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08805
 40 SEQUENCE DESCRIPTION:
 GATCCCGAAT CCCAGAAATA ATTCATAGNG CACCTATCAT GACTATTTAT TTAATCTCAG 60
 AAGCTGAATC TGTACGTATT TTTCTCCAGG TCATCAGCTT TGGGNNNNN 109

45 SEQ ID NO:7561
 SEQUENCE LENGTH:40
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08807
 50 SEQUENCE DESCRIPTION:
 GATCCTACTC TCTTATTAAA ATCTTGAGTT TATGTTCAAA 40

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SEQ ID NO:7562

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08808

SEQUENCE DESCRIPTION:

GATCTGCCGA CCTCGGCCTC CCAAAGTGCT AGGATTACAG GTGTGAGCCA CCGCGCCCAG 60
 CCATAACTTT NCTCTTTTCT CCATTAGCTA AAAGATAATA GTAAAACGTA TATTTGAACA 120
 CTTAATTTTC TCTATATATC TTTTNAGGAT TATATTTAGT GTATTTCCCA ATTTTCCTAA 180
 AATGATTATC ACTTCTGTAA AGAAGAAAAC AATACTACTG GTATTGCATT TGGAAATATT 240
 TTNCTATTAA NNTGTCTCAT TTCCTTTNCA TGTATGCACA TACATGTNNT TNGTATAGCT 300
 CCTTTNNTCA AAANNTATTC TGNGGTTATT ATTTANNN 338

SEQ ID NO:7563

SEQUENCE LENGTH:338

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08809

SEQUENCE DESCRIPTION:

GATCTTGTGG TCAAGAGCAA TCTGAATCCA AATGCAAAGN AGTTTGTCCC TGGGGTGAAG 60
 TACGGAAATA TTAAAGTAGA CGGGGCCCTC TTTTGGTGGA TGTAGCACA TTNCCACACT 120
 GTGAAGGCAG TATTAGAAGA CTTAATTGTA AAANCNCTCT TGTCACGTG TTACACTTAT 180
 GCATTGCCAA AGTTTTNTN AGTCTTGCAT GCTTAATAAA NGTNCTGAGA CTGTTACTAA 240
 GTAAAANGCT GTCAANCATT TCCTGANAAT AGANTTGGCC CCATGGCTTG ATGTGAAGAC 300
 AGCAAGGNAA GANGCACCAG TCAAGTTGTG ANCAAGNN 338

SEQ ID NO:7564

SEQUENCE LENGTH:324

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08810

SEQUENCE DESCRIPTION:

GATCCCTGCC ACAAGTTCTT TGGGCAGTGG CCATGTCACC ATTGAGATGA AGATATACAA 60
 CAGAAAATAG TGGCTGTGTT TGGAAGCTTC AGCCCTGCAC ATTTGAACTA GTCACCTCTC 120
 CAGACTTGCG TGGGTCAGTT CTTTCTGAGT AGAGGACTTG CTGGTAAAGG GGCAGATGCT 180
 TTTTATTAGT ACTGATAAAA CAACTGAGG GAAACATCCC TCTTAGCTGG GAAACTTTTA 240
 CTCTTCAGGA GCTTGGCATC ATGGACTGTT AATGTATGTG ATTTTCCCCC TATTTTCTCT 300
 CTCCAAAATG ATAAAAACAA TAAN 324

SEQ ID NO:7565

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08811

SEQUENCE DESCRIPTION:

5 GATCCCACTA CGAAACGCTC CAGAGGCTTC GGTTCGTCA CGTTCGCAGA CCCAGCAAAGT 60
 GTAGATAAAG TATTAGGTCA GCCCCACCAT GAGTTAGATT CCAAGACGAT TGACCCCAAA 120
 GTTGCAATTC CTCGTCGAGC GCAACCCAAG GTGTTCTAGC TGAGCGTTTA AGAGCATGGA 180
 CTCTGGAACC AGACTTTGAA TCCTTGCTCT GCCACTGCAG CTGTGTGACC TTGAGCAAAGC 240
 TATCTAAATA TTCTGTGCCT TTGTTTGTG ATCTGTACAA TGGAGATGAT GATGGTATCA 300
 CCTTCATGGG GTTGTCA 318

10 SEQ ID NO:7566
 SEQUENCE LENGTH:313
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08812

15 SEQUENCE DESCRIPTION:
 GATCTGAGAC TCAGGAACCT TGGCCCCCTT GTCACCTACA ACTGTGTGTG AACAGGACTA 60
 ACAGTTTGGA CTGAGAGCAC TGANGCTGCT GTGTGTTGTG CATATAACTT AATGCTGTGT 120
 GTTGTGGCTT GCACTGCTGT GGGGGCGGGG AGGGCAGTGG TGGGGAATGG CTGGCATGAA 180
 GCTTGCTTGA GCCAATGCTA TGCTTGCTGG GCTTTGGCTG ACTGGCTTTG GGCCACCAAC 240
 20 TGTATTGCA TAGTGATGGC TTGGAAGTGT CCATAGAAGG GATTGCAATA AAGGTGTGTC 300
 TTCTTCTAA AAA 313

25 SEQ ID NO:7567
 SEQUENCE LENGTH:316
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08813

30 SEQUENCE DESCRIPTION:
 GATCTTGAGT CAGAGAGGAT TGGAAGTGCT TTCTCCTCCA CCCAGGTGAG GTCAGGGGAG 60
 CTTAGGTCTT AGGGAGATGG CAAGTTGAGG TATGAAGGGA AGCTGGGGCT TTTGAGCTG 120
 CCGAACAACT GAGGGACCCA GTGCGCCTTC CATCCCGCAC TAGTGAATAG CGCCCCCTCT 180
 TCCCCGAAA ACGAGGTGCG AGAGGAACAA TTCCCACGCT GGGGAAGGAC TTGTCTCCTT 240
 35 TTCTGTGAAA ATGCTTTGTA AAAAGTTGTT ATTGTTTGCA TAGAGCAGAT TCTTGAGAAA 300
 AACTGTTTGG GACCAN 316

40 SEQ ID NO:7568
 SEQUENCE LENGTH:316
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08814

45 SEQUENCE DESCRIPTION:
 GATCCCGGTC TGCTGACCAG CTAACACCA ATTTCTTCCT GCTTTTCTCC ACTTGGTTTG 60
 GAAATCACA CAGTTTTAG GCTCCATCTG TTTGGAGAAA ATACATTCTG AAGCATCCCC 120
 AATTCACCTT CTA AAAAATC ATGTGCAGGT TTGATAAACA CCAGAACAGA AGACAGTGAT 180
 GCTGTATTAT TTTAGATTTA TTACATAGAT TTGGAATTCA CTTTTTTCAT GACCTAGAAA 240
 AAAACATTCC AGTGTTTANC TGTTTTNTAT TATTANGGGG CTTTTNATTT GTGAACCTCT 300
 50 GANGGCATGA GTGTTN 316

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EP 0 679 716 A1

SEQ ID NO:7569
 SEQUENCE LENGTH:226
 SEQUENCE TYPE:nucleic acid
 5 TOPOLOGY:linear
 CLONE:HUMGS08815
 SEQUENCE DESCRIPTION:
 GATCTGGTGT ACCAGGCGAA GTGGTCCGAG CAGGCTGAGC GATACGACGA AATGGTGGAG 60
 TCAATNAAGA AAGTAGCAGG GATGGATGTG GAGCTGACAG TTGAAGAAAG AAACCTCCTA 120
 10 TCTNTTGCAT ATAAGANTGT NATTTGGNGCT AGNNGAGCCT CCTGGAGAAT AATCAGCAGC 180
 ATTGANCAGA ANGAAGAAAN CAAGGGAGGA GAAGACAAGC TAAANN 226

SEQ ID NO:7570
 15 SEQUENCE LENGTH:306
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08816
 SEQUENCE DESCRIPTION:
 20 GATCAAGAGT TAAATAAAAA GCCCCAGGAC GAAACAAGAN GAGAGCACTT CATGCCCTGT 60
 GGTACAAAAA AGAGCAGCTC TACTTAATGC AATTCTTCAT TCAGCATATA GCGAGACATT 120
 TCTCCTGCCT CATTGAAAG ACATCCCAGC ACAGCATATC ACGCTGTTTC TTAAGTATTT 180
 GTATNNNNNN GTACCTGAAG TGTAGCGAAA ATNCTACTAT GACTCTNCCT GGAATACACC 240
 25 CACCTACCTT GANCCAGATT ATGGATTGGN TATGTCTACT TCTGGATGCA AATTTTNCCTG 300
 TTGTTN 306

SEQ ID NO:7571
 30 SEQUENCE LENGTH:304
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08817
 SEQUENCE DESCRIPTION:
 35 GATCCTGCGA GGGTGTCTATA ATACCTTGCT TTTCCATGCT TCCTGTATAT TTATCCATTT 60
 CAGCACATCT GCAGTAGCAA TAACTNCTNA TTTNTAAATT TNCTTTTGTT GGGGGAGACA 120
 GGCTTATGGC TGTCTGAGCA GCTGCTGTAG TGTAAGCAGT GAGGTTAGCA GTGGTTGTNA 180
 GATTNCTCA TTCCTGAGCA CCATGCTGTT GTGTCAGCAG TTGTAATGGA CTGTGTGGGC 240
 40 ATACTTCCAG TAGGTGATGC TTGCAAGTAN GANCCAGCTG CAGTGGTAGC AGTAGGGTTT 300
 GTAN 304

SEQ ID NO:7572
 45 SEQUENCE LENGTH:30
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08818
 SEQUENCE DESCRIPTION:
 50 GATCTGTACA AATAAAATNC CAAGTCCAAA 30

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EP 0 679 716 A1

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SEQ ID NO:7573
SEQUENCE LENGTH:118
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08819
SEQUENCE DESCRIPTION:
GATCATCTCT CCTTGGGTTT TNNTTTAAAA AGGGGAATCT NCTATAAAGG TTCTNTTGCT 60
TCAAACCAAT GTCAAATAGA CTTNATTTTA AGAGTCATGG AATTACAGTG CAACCTTN 118

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SEQ ID NO:7574
SEQUENCE LENGTH:53
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08820
SEQUENCE DESCRIPTION:
GATCTACCTA TACAGTCCTA CATTAGCTTC TAAAATATTT GTCAGGAGGG AAA 53

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SEQ ID NO:7575
SEQUENCE LENGTH:102
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08821
SEQUENCE DESCRIPTION:
GATCCCGAGG CTTTGTCTTC CTCTCGTCAG TTCTTTTGGT TGTGTTTTTT GTTTTNTTT 60
NAANAACCTCA AAAAAAAAAAN AAAAGACTTG GAGGAAGGGA AA 102

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SEQ ID NO:7576
SEQUENCE LENGTH:362
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08822
SEQUENCE DESCRIPTION:
GATCCTACTG ACAGCCATCG TGCAGGTGCT CAGCTGCTTC TNTCTCTATG TCTGGTCCTT 60
CTGGCTTCTG GCTCCAGGCC GGGCCCTTNA CCTCCTGTGG GTGAATGTNC TGGGCCCTG 120
GTTCAGTGCA GACAGTGGCA CCCCAGCACC AGAGCACAAT GAGAAACGGC AGCGCCGACA 180
GGAGCGGCGG CAGATGAAGC GGTATAGCC ATTGANATTG TGGCCACAGG CCACTGGGCC 240
CTGGGTGGCT CTGTNAGGGT GCACAGCCCC TCATGCCTGG AGCAATGAGG GTCTAGTCCA 300
GGGGCCAAAA GCAGTCTTGA GGGTATTGGG GTATACTTAT ACTCTATAGG GGTCGTTTGA 360
AN 362

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SEQ ID NO:7577
SEQUENCE LENGTH:109
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

EP 0 679 716 A1

CLONE:HUMGS08823

SEQUENCE DESCRIPTION:

5 GATCACTCAT ATAACATGTC ATTTTTTTGG TTCTGTTTTG GTTTGGTTTT TGCCAATNAT 60
TTTGTATAT TTCCAAAAA CTAATAAAA ATNATTTTAA TTTTNNAAA 109

SEQ ID NO:7578

SEQUENCE LENGTH:361

10 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08824

SEQUENCE DESCRIPTION:

15 GATCTTCACG GGGCTATTAT TTCAGTGACA AAATCCAAAT GCCCTCTTA TGTGGGTATT 60
ACAGGAATCC TTCTACAGGA AACAAAGCAC ATTTTCAAAA TTATCACCAA AGAAGACCGC 120
CTGAAAGTTA TCCCAAGCT AAAGTGGTG TTCAGTGTGG AAACCGATGG CTTTATTTCC 180
TACATTTACG GGAGCAAATT CCAGCTTCGG TCAAGTGAAC GGTCTGCGAA GAAGTTCAAA 240
GCGAAGGGNN CGATTGACCT GTGANTTCTT TGCCGTCTAA GGCAGTTGTT TATGACAGCT 300
20 GAAACTGGA CACTCCCTAA NTGTCCACCT TTCAGTGAAG NGATAGTTAA GCCAATTCN 360
N 361

SEQ ID NO:7579

SEQUENCE LENGTH:323

25 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08825

SEQUENCE DESCRIPTION:

30 GATCCAATTC GTGTCTGTGG ATGCTGCAAG GACTGGCTCA TTGGAGCCAA ACCATGTGGC 60
GGAAACTCCT GAAATTGGTG GAAAAAGTGG AATATGGCAG ATTGGGTAGG TAGGAGAACA 120
ATTTTGGAGA TTTTTTCT TATGTAAAG TGTCTGCATT TTATTGCTG TACCCTGCCT 180
ACTCTGGTTT TNAGGATTTT TTGACTTCTT CGAAAAGCAC TGAATAATCT CTTTNCCGGG 240
TCTAGATTTT TTGATGAGTA CTGTGTAATA TTGTCAATAT TCAATGGATT ATTTAATAAT 300
35 ANNTTAAGAN TTCCACTCAT AAN 323

SEQ ID NO:7580

SEQUENCE LENGTH:183

40 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08826

SEQUENCE DESCRIPTION:

45 GATCCTAAGT TAAGACTCTC CAAAGCAAAC ANTTGGTTTA AATCATTTAT CTGATGTAGA 60
TAAAAGGTAG AGAACTTGT AGTGTCTCC ACATAATGGA AAATGTACAA ATGCCTAGTT 120
GTGTTGCCCA TCAATTTTGT ATATNTCCAT AATTTTNATT GTTGANAATT GCATTATNTT 180
TTN 183

SEQ ID NO:7581

50 SEQUENCE LENGTH:350

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08827

SEQUENCE DESCRIPTION:

GATCCACCTG CCTCGGCCTC CCAAAGTGCT GGGATTACAG GCAGGAGCCA CCGCGCCTGG 60
GCCTTTTTTT AAGTTTTTAA NNCCNATAAA GNCACTGAAA GGTGATGTGT GTGGATGAGC 120
TAGGAAGACC TGAAATAGGC TCTCTCTAAA TTAATCAAAT TAATCCTGAA GCCATTCTGC 180
AATACTGTCT TTAATGTATA CTCATTGTGT ATAGAAGCCA GGGTTTTTNC CCCTAATTTG 240
TATCATTGCT ATATGTGTTA TTGTACCAAA CTACACTGTT TTAATTGCTG TAAATTTTAA 300
TATGCTTAG TATCTGGGTG TGGGAATCTT GANAGCATGG AGTTTGTGTN 350

SEQ ID NO:7582

SEQUENCE LENGTH:348

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08828

SEQUENCE DESCRIPTION:

GATCCTTGGG AGAGCTATCA CGGGAATCCA GGCCTTCACG GCCAGCAAGT GAGCCGCTCC 60
ATCAGGGGCC CGAAACTNTC AAGCCCCCTT NTGGAGAGNT GCCTGGCTGG ACCGTAGAGC 120
GCTGTGCTCT GAGCTTAGAA AGGGAGGTGG CGGATGGAGT GGGAAAGTGAG AGACACTGAT 180
TTTNAAATAT CAAAATTTCC CTTCTGAAGT CGTTCAGATG TTTTCCTTAA AAAGAAGATG 240
GAATTCTCTG TAGAGCGTCT CANTCCACTT TTAACCATGG NTGAGAGCAG ACTCCATTTA 300
CCNTGAAATA GCAGCTTCTT TTGAGGGGAG AGTGACATGG AAGCAACN 348

SEQ ID NO:7583

SEQUENCE LENGTH:258

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08829

SEQUENCE DESCRIPTION:

GATCTGCAGG TTTGGGCATA TGCTTTCATC ATTAAATTAT CTGATAAAGT TACAAGTCAC 60
AAAGGAGAAT GAGAACTTAA TGATTCTATT GGATTTAATA TATTAGCAAG AAANCATACT 120
ATTTNCATAT GTNTAGCTTA GTAAGGCATT ANCATAAGTN CAANACCTAT GANACAGATG 180
CNTATTTCTT CAACATACTG TGTCAGGTAT ACTGTTTTAT ANTTNGGTTG TTTTAGCCTT 240
ATTGCACACC NNCTCCCN 258

SEQ ID NO:7584

SEQUENCE LENGTH:308

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08830

SEQUENCE DESCRIPTION:

GATCCATTTT ATTACAAGTA AAATATTACA AAATATTTAC AGGAAAATAT TAAAAATAAC 60
TCAAACACAA AAGGCAAGAT GAAATAAACT GTTTTTTTTT CTCCAGAAAT CTCTACTCCA 120
GTGCCACAG CACACAAGAG TCAAACAAA TAAGCAACTA AGNTCCTTAC GGAAAAGGNA 180

EP 0 679 716 A1

CAGNTTGTTC CTAACCAGA AGAGGAGGTT GCCCAGAAGA AAAAGATATC CCGGAGGAAC 240
CTGAGGANCC NAACTTTTT GCCCGGGGG TAATTCNGNC TTTNNATTAA TTTNTTTAN 300
NGGGNNNN 308

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SEQ ID NO:7585
SEQUENCE LENGTH:36
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08831
SEQUENCE DESCRIPTION:
GATCTCTTAT CATAATAAG AATCTTTGTC ATCAA 36

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SEQ ID NO:7586
SEQUENCE LENGTH:227
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08832
SEQUENCE DESCRIPTION:
GATCTAAGGA GGCCAGTGAA TTTCAAACAG GATTAAAAAG AAAAAAACA GAGTTAGGTA 60
TCAAATTAAG AATACTGAAA GCAAAGATAT AANTAGAACT TAAAAGCAG CCAGAAAAAC 120
AAACATATTA CATATAGGGC AGCAACANTA AGANTAACCG CTAACTTTG ATTAANNNGTA 180
ATATAATTCA CAATACANTG TAACANCANT TTTNNGTGCT AACNGGN 227

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SEQ ID NO:7587
SEQUENCE LENGTH:215
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08833
SEQUENCE DESCRIPTION:
GATCTATTGA GAATGGAATT GTTTCTTAA TTTCATTTTC AAATTGTTTG TTGCTAGCAT 60
ATAGAAATAC AAGTGATTTT NTATATTGCT CTTGTATCCT GTGACTTTGC TGAGCTTATT 120
TTAGTAGTTT TTTTCACTG AATTCCTTAG GATTTNTAC ATAAAAAATT CATATCATCT 180
GTGAATAAAG ACAGTTTTAT TTCTNAATT TAAA 215

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SEQ ID NO:7588
SEQUENCE LENGTH:43
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08834
SEQUENCE DESCRIPTION:
GATCATATTT TAAATAAAG CATTTTTTGT AGAAAGTGCT AAA 43

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SEQ ID NO:7589
SEQUENCE LENGTH:334
SEQUENCE TYPE:nucleic acid

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TOPOLOGY:linear

CLONE:HUMGS08835

SEQUENCE DESCRIPTION:

5 GATCAGGATA ACCAGGATGA TAGCAGTGAC GATGGATTCC TCGCTGATGA CAACTGCAGT 60
 TCAGAAATAG GACAGTGGCA TNTAAAGCCT ACTATCTGTA AACAACCTTG TATCCTACTT 120
 ATGGNCTCAC TCCGAGGCCC TTCTCGGTCA AATGTTGTCA AAATTTAAG AGAGTATTTA 180
 GANGTGGAAT GGGAAAGTTAA AANAGGAAGC NNNAGANGTT TTTCCAAAGA GAAAAGAAAG 240
 10 CATAAGGNCA CTTACTCAAC AGANGCACCT TTAGGCGGNG GANCAGNACA TTGTGTCAAT 300
 AGTNTCTCNG GTTTGCCCTT TCTNTTACTT GTCN 334

SEQ ID NO:7590

SEQUENCE LENGTH:235

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08836

SEQUENCE DESCRIPTION:

20 GATCAAAATTT AAAAATCAGT TTTAAAAGTG GTTCCCGACT TCTTTCCTTT GGAAGAATTT 60
 GGNCTNATGC TCATAAGCAG TTCATGAGTG ACAAGTGTAG CAAAAGCCAG CCATATATTT 120
 CNAGANTCTA TATCCTCAGG AAATGGTCCT TTTTTCCT ATAANTCACC AACCAACTAT 180
 CTAANTGGNC ATTTGGGGGA NTTCCCTCCC AAATTGATT TTNTNATNG AAAAN 235

25 SEQ ID NO:7591

SEQUENCE LENGTH:172

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08837

30 SEQUENCE DESCRIPTION:

GATCCAGAGC ACAGATGGCA AGITGTGTGG ATGATACTGA ATATNTTAT TTTCTAAAG 60
 ATAAATNTCA CGACACGACA TTTCTCAGTT TTAATTTAGA TGTTATNAC CATGTGTAA 120
 CAGAATCTTA AAACCAAGG NATTTCTTCA GTAAACTAGA CTTTGATTCA AA 172

35 SEQ ID NO:7592

SEQUENCE LENGTH:335

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

40 CLONE:HUMGS08838

SEQUENCE DESCRIPTION:

45 GATCAAGAGG AAGCATGACT TGCACAAGAT GGCAGAGGCC AACCGTGCCC TGGCCCACTA 60
 CCGCTGGTGG TAGAGTCTCC AGGAGGAGCC CAGGGCCCTC TGCCGCAAGA AACAGTGTGA 120
 GCTACTGCCA CGCTGAAAAC TACCTGTGGG TTAAGGATGT AGTTCCTTTG TAAGGGTGGG 180
 CAGGCCTCGT AAGAAAGATG TAGCAGCATA TTCACTATCC GTTAATCCTT CTTTCTTTGA 240
 GGCTGGAACT TGCTCTCTCT GCCCCTATTT CTTGTAAAG AGGGAGCACA TTGACTTGGG 300
 AATTTCTTCC AGGAAACTCA GGGCTGTTTT CTCTN 335

50 SEQ ID NO:7593

55

SEQUENCE LENGTH:333

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08839

SEQUENCE DESCRIPTION:

GATCTGGGAC TTCTTTAGTA TCGAGTCTCC TTTGCACTCT TAAAAATTGT TGAGGACCCC 60
 AGAGAGCTTT TGTTTTATG GGTGTATCT ATCAATATT ATCACATGAG AAATNATTTA 120
 TTTATTTTAA AATAACTTTT TTATTGAGAA AGATAGCAGT GTTTTTTTTT AATCTTGCAA 180
 ATTGTTTAAT GTTTGATTGA ATGGAAGACA GCCAGATTGT CCTATCAGCA TTCAAAGTGC 240
 TGCAGTAAAGT TGTTTGACT AATGTATATG AGGAAAACCT GACCTCACTG AGATTTTAAT 300
 TGGAAAATGG AGGAGTATTT TNCCAGCCTT TNN 333

SEQ ID NO:7594

SEQUENCE LENGTH:220

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08840

SEQUENCE DESCRIPTION:

GATCTCCTCC ATACGGAAC TTCTCATCTA CGTGGCCCTC CTGCGAGTCA CTCCATTTAT 60
 CTTAAAGAAA TTGGACAGCA TATGAAGACA GGACATCACA TATGAATGCA CGATATGAAG 120
 AGCCTGGTTA CAGTTTCGAC TCCTCTCTGC AAGTGAATAG GCCCAGAAAG GTGTAAGAGA 180
 CTCTTTGAAT GGACATAAAA TTCTGCTTGT TAAGAACAAA 220

SEQ ID NO:7595

SEQUENCE LENGTH:326

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08841

SEQUENCE DESCRIPTION:

GATCTGTGCT GGCCACGTGC CCGGNGTCTC AGCTCTGAAC CTGCTGTCCC TGCTGAGAAG 60
 CTCTGAGGGC CCCTCCCTGG AGGACCTGTG AGGGTGGCTG GCCCCTGGGC TGCCCTTCT 120
 NATGGCTTCG TGCTGACTCC ATAAACATTC TCTGTTGAGG ATGTCCAGTC AGGGCTTGAC 180
 AGGCCAGGN TCAGCCCGCC GTGGCTGGGA AGGTTCCCTG CAGTGCCAGT GCTGCAGCAG 240
 GGAGAGCTGG GCAGAAGCAG CGAGGGGGCC CAGCTGGCGA GACTGTAGCC CCGTCCCACT 300
 CCCACACTCA CTCTTGTTAG AGCCTN 326

SEQ ID NO:7596

SEQUENCE LENGTH:327

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08842

SEQUENCE DESCRIPTION:

GATCAGATGT TGAACGTGC TATGCTGTAG CGTGTCTGGA AACAAAGTAC ACAAACCTGG 60
 CTACGGTGAT GAGTTAGCTT CTGCTTACTA CCTGTNACAA CCCAAGTGGG TGACACTAGT 120
 GAACCTTCTC CAGTCTGCAG GCTGGCATAG AAGGCTCTTA GATTATATTG GGCAGCTTGC 180

EP 0 679 716 A1

AATCTGCCGA AGCAGTGACT TGCATTTCCT CACTTGGCTT GAGCACTCAA CCCAGAAGGC 240
 GAAGATAGCT TTTGGTTGTA GCGGGCTTCC TGTATGGGAT ATCCCTCGGT AAGGGTAAAG 300
 GAGCAGAGGC AAAGGAGAAA AGCAGTN 327

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SEQ ID NO:7597
 SEQUENCE LENGTH:278
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS08843

SEQUENCE DESCRIPTION:

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GATCTTTCCT CCCTCCCCTG CATGAGGCAG AGGCAAGCTG CCTGCCAACC CCCTCCCTCA 60
 AGGAATGGCC TTGCCAGGA ATGCCACCA CACATACCCT CTNCTTTTT NNTAGTCAA 120
 CTCTTGTTA TTCCTTGGNT TGCNTCCCTC CTCCTCCCC TCTCAACCTT TACTTCTGAT 180
 TTNNAATTCA TGGAAATTGG GATTGANGTT AACTACAAC AGTGCCGCCA ACACCAAGTN 240
 TTGCAGGAAA AAAATACAAA GAAATTTANC ANANAANN 278

20

SEQ ID NO:7598
 SEQUENCE LENGTH:202
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08844

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SEQUENCE DESCRIPTION:

30

GATCTTCATT ACCGTTNCAT GATTGGAAAT ANTAAAACA TTGTACAGTT TTAGTATAGA 60
 GAAATNTAAT GGNTTTTGT AACAGTTTC TGTCTGCATG TAATTGGAT TTCTCAAATA 120
 CATTCAATAG TAATTNATCA GTAACATTAG TTTTATTNN GTCCATCTCC TTATCTATAA 180
 AANGGGGATA TNCTTAGGAT NN 202

35

SEQ ID NO:7599
 SEQUENCE LENGTH:47
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08845
 SEQUENCE DESCRIPTION:
 GATCCATAAT ATAGCATCAC ATCTCAATAA ATTTATTTGG AAGCAAA 47

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SEQ ID NO:7600
 SEQUENCE LENGTH:327
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08846

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SEQUENCE DESCRIPTION:

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GATCTGCCCA CCTTGGCCTC CCAAAGTGCT GGGATTACAG GCATGAGCCA CTGTGCCTGG 60
 CCCCTTCCTG TAAAAATTTN AAATGGAGAA TTGGGTGCNA GATGTGGTTT CCAGCCTGGT 120
 GCCTGGGGTG CTGAGCTANN NGAGTGGTGC AGTCCAGGAC ACCTTTGCTT TATCTCACTT 180
 ACACGGTCAC CTGGAGCCGG CTCAAGTGGC TAAAGCATCC TGGGGCCAG AGCCAGGTGA 240

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EP 0 679 716 A1

TAGGTCCCTN TGGCCAACTG GACAGTTGAG GCCTGTGGTT ACCCGAAGCC CAGCTTGGGG 300
CCCTGGTCCA GCCTNGCCTC CCAGGAN 327

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SEQ ID NO:7601
SEQUENCE LENGTH:326
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08848

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SEQUENCE DESCRIPTION:
GATCTAGATG CTGGTATTGA GGTGGGAGAC AAGTACTGCC ACCNGAAACA ACAGTCTTGG 60
TAAATTTAGC CGACGAGGGT AAACACATCC TAACAGGGAA GGTAAACTGT ACGTCCATCA 120
GTACCACTAG AGGGCATCNC NGGTTTATAG TTCAATACAG TGAATATATC AGANTAATGG 180
CCTTTAGTTT TCCTGAAAGA TTAANTTAGG CTTGCTAACT TGTTTAATGN GATANTCAAA 240
CATATGATGT AATTTTAAAG GGTTTACATT TTTANAANTT TAATGGTTGC TACATAAAAN 300
GGGNTTCAGT TAACTNATTT TGC GTN 326

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SEQ ID NO:7602
SEQUENCE LENGTH:310
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08849

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SEQUENCE DESCRIPTION:
GATCTCAGCA GGGAAACCATG GAAGCACCCA CCCCTTGCA TTTTGGCTTA CCTGTAGCGG 60
AATGCAACAG AGAATTTTAT ATGCTAAGAT TTGTGGTTCC GTGGGAGAAT ATTTCAAACA 120
GGTGGAACC ATCAAATAA AAGAGAACCA GAAATCATTT AAAGGTCACG TGAGGAGCTT 180
CTGTNTTGTG CAACTTAAGC CTTACCAGT CTCCTATGCC CAGGTGCCCA GCACCCCAT 240
CATCTGCATA TTCACCATCC TCAGGTCAGA TTCTATGATA CTGNNATAA AGGCATTCTG 300
GAAAAATAAA 310

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SEQ ID NO:7603
SEQUENCE LENGTH:287
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08850

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SEQUENCE DESCRIPTION:
GATCCGGTCA GCAGCCCTG TGGCTAGAGG AACACCAGCA CAAACGACAG CCTCAAGTCT 60
CCTTCGAGCT TTATATCCAT TTGGGGATGA AGTCTACTTT NACAGCTAGC AAGGCGACAT 120
GCAACTGTTG TTGAATGATG ACAGCAATTC AGGAAAGACT TAAATATGAA AGCAAATTGA 180
NCACATCGGG TGTGTTGNAT CAGANNAGNG NTGNGATGTG ATAAGACTTG TTTATTGACT 240
AGCCAATATG TCATTACCT TCNGGTTTAT NTTGTGNANC CCCANN 287

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SEQ ID NO:7604
SEQUENCE LENGTH:66
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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EP 0 679 716 A1

CLONE:HUMGS08851

SEQUENCE DESCRIPTION:

GATCGAGTTT TGGTTCGGGT TAACTGTGTG CCTACTGAAC CTGGCAAATA AACATCACCC 60
TGCAAA 66

SEQ ID NO:7605

SEQUENCE LENGTH:339

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08852

SEQUENCE DESCRIPTION:

GATCTGGAAT GTGAAGCGTT ATAGAAGATA ACTGGCCTCA TTTCTTCAAA ATATCAAGTG 60
TTGGGAAAGA AAAAAGGAAG TGGAAATGGGT AACTCTTCTT GATTAAAAGT TATGTAATAA 120
CCAAATGCAA TGTGAAATAT TTTACTGGAC TCTATTTTGA AAAACCATCT GTAAAAGACT 180
GAGGTGGGGG TGGGAGGCCA GCACGGTGGT GAGGCAGTTG AGAAAATTG ANTGTGGATT 240
AGATTTTGAN TGATATTGGA TAATTATTGG TAATTTTATG AGCTGTGAGA AGGGTGTGTG 300
AGTTTATAAA AGACTGTCTT AATTGCATA CTTAAGNTN 339

SEQ ID NO:7606

SEQUENCE LENGTH:337

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08853

SEQUENCE DESCRIPTION:

GATCCTTGTG GAGCAAGTTC TATTGTTC TTTGCTTGCCA GAGATGAACA CAGAATGTTC 60
TGTTTCATTT TACAAGAACT ATCCTGAGTT TCTGTGGATG GAAACATTAC ATGTAATGCA 120
GATATAGTGA AACTTGGAAT GATTATTAA AGANTTATAT TTGTGTATAC TTTATAAATT 180
AGTCCCTCAT TAGATTTTTT TTTTCTTAAG CATAAGACTG ANCTTAAATG TGTTAATTTT 240
AGTAGAATCA GGCAGTCTC GCAGAAGGAN CACAGATTGT NGAGATTAAC ATAAATNGTT 300
CTTGTCTCAA TATATATNTN TNNCCATTTC TGTCGAN 337

SEQ ID NO:7607

SEQUENCE LENGTH:203

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08854

SEQUENCE DESCRIPTION:

GATCCGTGGA TTGAGGGTGC CTTCCATTT ATTTGGTCT TCTTTACTTT CTTCAACCA 60
AGTTTTGTAG TTTTCAGAGT ATAAATTTA TACTTGTTTT GTTCAGCCTA TTAGTGTTTT 120
ATTTTGGTG CTATTGTAAA TNAATCTTC TTAATTTCAA TTTTCAGATTA TTAGTTAAA 180
TATACAGAAA TACAGCTGAT AAA 203

SEQ ID NO:7608

SEQUENCE LENGTH:207

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear
CLONE:HUMGS08855

SEQUENCE DESCRIPTION:

5 GATCAAACT TTTTATAATA CAAAGACATT ATTATTTGNT TTTTCACTG TCCCCTCCTG 60
AATATACAAC GGAGCATTCC AGATGCTAGA GTGACAGCCT CACTCAGACT CAGGGGATGC 120
ATGCTTG CNT GTACAGTACT CGTATGTTT AAATTGTTT TAACTCCTAA TATGTAAGTA 180
TCAATAAATA TAATCACCTT AAGTAAA 207

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SEQ ID NO:7609
SEQUENCE LENGTH:249
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS08856

SEQUENCE DESCRIPTION:

20 GATCTTCGTG GTGGAAAGCT AAATTTTAAA ACCACCCCAA TGGATGCAGA CAGTGATGTT 60
GCATTGGACA TTCTAATTAC AAATGTAGTC TGTGTTTNA GAACAAGATG TCATTTAANC 120
TTAAGGAAGA TTGCTTTGGA AGGNGCAAAT GTAATTNATA ANCGTGATGT TGGAAAAGTA 180
TTANTGAGGC TTAGAAAACC TAGANTTACA GCTACANTTN GGTCTCAGG AAANNTTATT 240
TGCNCTGGN 249

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SEQ ID NO:7610
SEQUENCE LENGTH:319
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08857

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SEQUENCE DESCRIPTION:

35 GATCTGTAGT ATTTAATGT GCATCTACTT TAAATGAGTC ATCTGGGGT TTTTATAATT 60
CCCTTATGTT CTCGCCCTC TACACTTGAA ATAACAAAAT GCCTTAATTT TATGGATTAG 120
NTCTCTTATA GGAGACAGGC AGCTATATGN AGCAANCCCA GTAAAGTTAT TTTTCAACTN 180
TCANAGTTGT AAAATATTTT ATAACAGTTT ACANCAACAG CTGNAGANCA ACATGCCACA 240
TTTTNTTGC AGCATTTTCA AATANTTTGT TTTGTGGTGT NAGCACAGGN TAAANNTGGG 300
GNGCGTCGAT GTGAGGCGN 319

40

SEQ ID NO:7611
SEQUENCE LENGTH:208
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08858

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SEQUENCE DESCRIPTION:

50 GATCTACAAA AAATTCCTGG AGCCATATAT ATACCCTCTG GTTCCCCCT TCGTTAGTCG 60
TATATGGCCT ANGAAAGCAA TACAAGANTC CAATGATACA AACAANGGCA ANGTAACCTT 120
TAAGGGTGCA GACATGAATG GTTTACCAAC AAAAGGNCCA ACAGAAATCT GTGATAANNC 180
GNAAGACTAA NGAAATTTTC CNAAGGN 208

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SEQ ID NO:7612

55

SEQUENCE LENGTH:321
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08859
SEQUENCE DESCRIPTION:
GATCTTCCCA CCTCAGCCT TCCAAGTAGC TGGGACTACA GATGCACACC TCCAAACCTG 60
GGTAGTTTTT NAAGTTTTTT TGTAGAGGTG GTCTAGCCAT GTTGCCTAGG CTCCCGAACT 120
CCTGAGCTCA AGCAATCCTG CTTCAGCCTC CCAAAGTACT GGGATTACAG GCATCTNCTG 180
TAGTATATAG GTCATGAGGG ATATGGGATG TGGTACTTAT GAGACAGAAA TGCTTACAGG 240
ATGTTTTTNT GTAACCATCC TGGTCANCTT AGCAGAAATG CTGCGCTGGG TATAATAAAG 300
CTTTTCTACT NCTAGTCTAA A 321

SEQ ID NO:7613
SEQUENCE LENGTH:136
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08860
SEQUENCE DESCRIPTION:
GATCCTTCCT AGGGGATGGG GGAAGCCCTG GCTGCAGGCA GCCTTCCAGG CAATATGNNG 60
ATAGGAGGCC CACGGGCCTG GCAGTGAGAG GTGTGCCCC ACACCGATT ATGATATTAA 120
AATCTCAACT CCCAAA 136

SEQ ID NO:7614
SEQUENCE LENGTH:47
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08861
SEQUENCE DESCRIPTION:
GATCCCGTCT CTACAAAAA TAAAAAAA TTAGTCGTGG TGCCAAA 47

SEQ ID NO:7615
SEQUENCE LENGTH:36
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08862
SEQUENCE DESCRIPTION:
GATCACTTTT TCAATATTAA ATNTTATTTA CATAAA 36

SEQ ID NO:7616
SEQUENCE LENGTH:260
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08863
SEQUENCE DESCRIPTION:
GATCATCATG AAATNATAAG AGGGCTTAAG AATTTGTCCA TTTGCATTG GAAAAGAATG 60

EP 0 679 716 A1

5 ACCAGCAAAA GGTTACTAA TACCTCTCCC TTGGGGATT TAATGTCTGG TGCTGCCGCC 120
 TGAGTTTCAA GAATTAAAGC TGCAAGAGGA CTCCAGGAGC AAAAGAAACA CAATATAGAG 180
 GGTGGAGTT GTTAGCAATT TCATTCAAAA TGCCAAGTGG AGAAGTCTGT TTTAAATAC 240
 ATTTTGTGT TATTTTAAA 260

SEQ ID NO:7617
 SEQUENCE LENGTH:102
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08864
 SEQUENCE DESCRIPTION:
 15 GATCAACTTT AGTCACCAAC CCAATGGAG GATTNTGGC TTGTGGNCC TTATATGCCT 60
 ATAGNTGTGG ACATTGTCAT TACACAACTG GANTCTTNC TN 102

SEQ ID NO:7618
 SEQUENCE LENGTH:228
 20 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08865
 SEQUENCE DESCRIPTION:
 25 GATCAAGGGG CCTCTCAGAA CCATGTTCCC CAGCCAGGTG AGGACCATT TAACTGGGAC 60
 CCAGGNCAAA ACCATGTGGG TGCACAAAGC CAGGCACTGC CAAGTGGAAC ATGAGGTTAT 120
 TTCCAAATCA TGGGAGCCAC CAGCAGGGAG AGGGCAGTAT GGAAAATCCC CTGGAGCCGG 180
 TCAACTNTTT GCTCATGGCT AGTGAAATAA AGTTGTTTGA GTACTAAA 228

30 SEQ ID NO:7619
 SEQUENCE LENGTH:113
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS08866
 SEQUENCE DESCRIPTION:
 GATCTCTTAC TCCCCAGTT TGAATGGTAA ATTTGAATGG TAAATCCCA TGAACATATA 60
 TGGAATATC TTTATCCTAC TTTNTCCAAT AAAGGCTGTT CTTAGCTTTT AAA 113

40 SEQ ID NO:7620
 SEQUENCE LENGTH:279
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 45 CLONE:HUMGS08867
 SEQUENCE DESCRIPTION:
 GATCACCTGN CAAATGTTTG AGGACGGAGC TGTGCAGTCA CATTATTGGG GATTCCACAG 60
 CTGGTGCTGC AGGCCCTGCG CCTCCAACCA GGACTTTNTT CTTAATGCTC TCGACACTTA 120
 GCTAAACAG ACTATATCCC GGCCAGCAG GCCCCAGCGC CGTTAGTCTC CAGCTGACTC 180
 50 TGTGGGTTGG TCTTAAAGCA AATTCTGTTT TGTGGACTGC CTGGCAATTT TTNAGCTAAC 240
 TGTAATGATA AAAAGGGAGT NTAATCTAT TCTGAATCN 279

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SEQ ID NO:7621

SEQUENCE LENGTH:198

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08868

SEQUENCE DESCRIPTION:

GATCAAGTTG TGGTGGATAA CGTGTITCCA AATTGCATCT TGTGCTGAA ACTTCCTGGA 60
 CTTGAGAAGT TACTTCATCA TGTGACAGAG GAAAAAGGTA ATCCAGAAAT AGACAACAAG 120
 AAATATTACA AGTNCAGCAA AGNGAAGACA TTAAAGTGGC TGGNAANNNN GGTAAATCAA 180
 NCTGTGGCAG CATTAAAN 198

SEQ ID NO:7622

SEQUENCE LENGTH:184

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08869

SEQUENCE DESCRIPTION:

GATCCAGAAA GTAGCAATAT AGGACTTTAA GGGGTATGTA TTAGTGTAAT TAATATATGT 60
 GCAGTCTAGG CTAATCATNT TAATTAAAAA CAAATAAGCA TACTTTTTTN CTTAAAAAT 120
 ATGCCCTCAA AATGGTGGGG GATAAANTGT AAGATATTCT TTAGCTGNNN TTATTNNCC 180
 GNCN 184

SEQ ID NO:7623

SEQUENCE LENGTH:212

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08870

SEQUENCE DESCRIPTION:

GATCCTAAGT CTCTACAAA AGCTTTGAAT ACTGTGAAAA TTTTTACAT TCCATTTCAT 60
 TTGTGTTGTT TTTTAACTG CATTTTACCA GATGTTTGN TGNTATCGCT TATGTTAATA 120
 GTAATTCCCG TACGTGTTCA TTTTATTTTC ATGCTTTTTC AGCCATGTAT CAATATTCAC 180
 TTGACTAAAA TCACTCAATT AATCAATGNA AA 212

SEQ ID NO:7624

SEQUENCE LENGTH:373

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08871

SEQUENCE DESCRIPTION:

GATCGCAGAC CGCGCTGCCC ACGACGAGTC CCCGGGGAAC AACGTGGTTG GACATCAGCG 60
 TGCCGCCAC CACGGACGGA CAGAAGTCTC TTAAAGAAAA TAGTTTAAAC AATTNGNTAA 120
 AAAATTTTCC GTCTTATTTT ATTTCTGTAA CAGTTGATAT CTGGCTGTCC TTTTATAAT 180
 GCAGAGTGAG AACTTTCCCT ACCGTGTTTG ATAAATGTNG TCCAGGTCTT ATTGCCAAGA 240
 ATGTGTTGTC CAAAATGCCG TTTAGTTTTT AAAGATGGAA CTCCACCCTT TTGCTTGGTT 300

EP 0 679 716 A1

TTANGTATGT ATGGAATGTT TATGGATAGG GACATAGTAG TAGCGGTGGT CAGACCATGG 360
GAANTGGGTG GNN 373

5 SEQ ID NO:7625
SEQUENCE LENGTH:74
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS08872
SEQUENCE DESCRIPTION:
GATCCTACAA GCATGTCTGC TGTAACCTA TGTATGTTTT CTGTAAATC ACATACCTAT 60
AAAATAAAAA GAAA 74

15 SEQ ID NO:7626
SEQUENCE LENGTH:53
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
20 CLONE:HUMGS08873
SEQUENCE DESCRIPTION:
GATCGTGAAC ACTGCATTCC AGCCTGGGTG ACAGAGCCNG ACCCTGCCTC AAA 53

25 SEQ ID NO:7627
SEQUENCE LENGTH:379
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
30 CLONE:HUMGS08874
SEQUENCE DESCRIPTION:
GATCTACCCA CCTTGGCCTC CCAAAGTGCT GGGATTACAG GCGTGAGCAC CGCGCCTGGC 60
CCTCTGCAGT TGTTTAATAA GGCACAGAAT ACCTGTAGCA TAGGTCAGCC TTACGATGTC 120
CATGAATTAC ATATTACAGC GTTTTAGAGC CTGATACATT TTGGAANCGA AAAACAACCTT 180
CTACACCTAT TCTACAGTCC GCATTTAAAA CAATAAATTC CTCTATTAAA AACGTAAAGC 240
35 CGGGTTTGCT TGCCTGCCAC AGGGGGTATA TCCAGGAAGG TTATTATGAA GCTGTCAAAT 300
CAAGATGATG GNAATAAGGC AGTTTGAACG GACAGTCTTC CCACAGTCAG GCCATTTTGT 360
CTGATTTGGT TTAGGATTN 379

40 SEQ ID NO:7628
SEQUENCE LENGTH:212
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
45 CLONE:HUMGS08875
SEQUENCE DESCRIPTION:
GATCGGGCGC TGTGCAGTGT GGTGAGCATG GTGAAGAAAG TCATTTCTC GGTGGGCAGT 60
ATTCTCTTT ATCTCTCATT ACACTGGAAA TGTATTCTT GCTGTATCAT CCGTGCTCAA 120
CGTTTTAGTC TGTGAGGCTC ACCTTCTCTC TGGAAAGAAT TTGCTTAACT TGACATTCCA 180
50 TGTGCCGCTA ATAAATATA TTTTGAAGA AA 212

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SEQ ID NO:7629

SEQUENCE LENGTH:350

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08876

SEQUENCE DESCRIPTION:

GATCAGAANT AATTACCATT CCCNGTGTTT TGCTGTAGGA CCTTCAGGTC TATGTTCTAA 60
 GTGGGGNTTA TGAAACTGGT AATAGAAAAT AATGGAGAGC AGTTAAAGAG TTAATATACA 120
 TTATTAATAT GNGTGCTCTC TACAGGCACA CAAAGCCTGT AGACTCAACA AATTTACTCC 180
 TGAAGTTTTC CAAATTTGCA AAATCTGCAA TTTNTGTGCC ACATAAGCT NTCCAGAAAT 240
 TGAAAANCNG TAAANCACCG GGTCTCANTT TTTCTCTGCA TTTCACTGAT ACCTTAANTT 300
 ACATAGNTTA NCTTCTTTAN AGCACTACCT TTTTNGCTGT TGNTATTAAN 350

SEQ ID NO:7630

SEQUENCE LENGTH:76

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08877

SEQUENCE DESCRIPTION:

GATCTCACAA CTGCACTCCA GCCTGGTGAC AGAGCAAGAC TCCATCAAAA ATAAACAAAT 60
 AAAACAAACA AACAAA 76

SEQ ID NO:7631

SEQUENCE LENGTH:185

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08878

SEQUENCE DESCRIPTION:

GATCAAAGTC TGCTTCATTG TCTTGTTCCC CATTTTAATG GCAATGTTAG GTATAATNTT 60
 TGCTTTTATT AAGTATAATN TTTGCTTTTG TTGGTAGAGT TTTACGATTT TACATCAGAA 120
 GTACCTGTTT AACTTAATTG CCACGTTTAA GCATCAATTA AAATATTTTC ATTGAACTCA 180
 TGAAA 185

SEQ ID NO:7632

SEQUENCE LENGTH:149

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08879

SEQUENCE DESCRIPTION:

GATCTTTTAT GAATTAACT TGCAAAATNAA ATGAAATATG TTGGGACTTT GGCATAATTT 60
 GAATATAGCA TTGTGTGAGT TGTGATTGTN TCTTCCTAAT TGTGCTGTAA TAACCTGTGT 120
 ATGAGAATAA AATGATTTC ACCATCAAA 149

SEQ ID NO:7633

SEQUENCE LENGTH:193

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08880
 SEQUENCE DESCRIPTION:
 GATCTCACTC CATGATTACT GTGTAAAATA TTTTGCAC TTTGTGAAGT ATTTTAACT 60
 TTTTCTGTA CATAACTGTG TTCTCAGAGC TGAATGTTA TATCTTTGC TGTGCAAAAG 120
 AAACANNNA AATGTTGTTT AGTTGTATAT ACAGAAATGT GTATAAACA TTTTGTATT 180
 TTTNAAAAGT AAA 193

SEQ ID NO:7634
 SEQUENCE LENGTH:148
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08881
 SEQUENCE DESCRIPTION:
 GATCAAGAAT TTTGTAGAGT GGACAGTCAT TACATATGTT ATAACCTATC CTTAAAAAC 60
 TATTTTAAAC TTTATCCTTT CAGCTTTACT TAGTGCGATG TTTTGAAGC AGTCTTCAAA 120
 GAATAAACA CTAACCATGC ATGTGAAA 148

SEQ ID NO:7635
 SEQUENCE LENGTH:376
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08882
 SEQUENCE DESCRIPTION:
 GATCCATTTT TGTGTCCTNT AGCCTAGTAT GTCTGAACTT CCATTCTTGT TATATATTTA 60
 AACTTTCCCT CTATATTATA GGNITNGTGG CNCCACGGT CAGGTGTAGA GGAAGCTGCC 120
 CCTTGCAGAA CTGACTGTA ATATTTTCT TTTATAAATA TTTTCACAGG ACTGATTGTA 180
 CACAGGGCTT GTAATAAAAT TTTAACTG TGCTGTGAAA CAACTATGGG GAATCTCCAT 240
 TGAAGGCTAC TTCATGGGCA CCTGAAAGTG GAGTGTATA GCTATGACTT TCTATTTCTT 300
 GTTTCCTAAG TAAATTAAAC CTAATTTTCA CCCTTTCATT CTGTTTCAGC CTCCTGTNTA 360
 AGGAAGTACC GTTGN 376

SEQ ID NO:7636
 SEQUENCE LENGTH:279
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08883
 SEQUENCE DESCRIPTION:
 GATCCCATTC ATTGTCCCCT TTGGGGTATT TCCAATACTT GAATGGCAGA TTGGAGTTTT 60
 TCAGAGTATG TGTTTCATCT GCTAGTCTTT CTCTCCTTCA TAGCTTTTCT TTTCTGGAC 120
 TTGCTCCTTT TGAGTTGCTT TTGCGTTTCT CATGCCTAGG CAAGTGTAAT AGAAATTATG 180
 TAGCTCCTTA TGTTGGCAA GAGCTCTAT ATAGTTTCAC TTTGTATACA AGTTAGGNCC 240
 AGCTGTTGTT ACATGTAATA TNNTGGTTCA GAACTTGAN 279

SEQ ID NO:7637

SEQUENCE LENGTH:373

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08884

SEQUENCE DESCRIPTION:

GATCTCTGAT TGCATCTATG TGGATTCTTT TTGGAGGTTA TGTGCTAAA GAAAAAGACA 60
 TAGTATACCC TGGAAATTGCT GTATTTTNCC AGAATGCCTT CATCTTTTTT GGAGGGCTGG 120
 TTTTAAAGTT TGGCCGCACT GAAGACTTAT GGCAGTGAAC ACATCTGATT TCCCACAGCA 180
 CAACAGCCCT GCATGGGTTT GTTTGTTTTT TTAGTGCTCA CTCCAACCT TTTGTAATGC 240
 CATTNCTAA ACTTATTCT GAGTGTAGTC TCAGCTTAAA GTTGTGTAAT ACTAAAAATCA 300
 CGAGAACC TAAACANCAN CCAAAAATCT ATTGTGGTAT GCACCTGATT AACTTATAAN 360
 ATGTTAGGTG GTN 373

SEQ ID NO:7638

SEQUENCE LENGTH:53

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08885

SEQUENCE DESCRIPTION:

GATCACACCA CTGGCACTCC AGCCTGGGCG ACAGAGTGAG ACTCCGTCTC AAA 53

SEQ ID NO:7639

SEQUENCE LENGTH:363

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08886

SEQUENCE DESCRIPTION:

GATCCGAAC TGGACTCACAT CCTGTATGGT GGATGGACTG TATATTGAGG GTTCCATTCT 60
 TCGCGCAGTT TAGACATCTC TGTTTGTATT CTTTGTGTT GTTTTATTT TAAAAGGCAC 120
 AAACCTCTAGA TATTAGTTGA ATGTTGAGGC TTAACTTTT TCGGTGTCTT TCTACAACTG 180
 TGTCTGTGA CTCAATTGTG TCGTGTTAAT ATCAGTACAG ACTGTCTCCT CTACGTGACC 240
 GTATAATGTT TTTCTCTTCT TGTAGTCTCT ATGGCGTGTC TTTATGGTGT AATAAGGTTT 300
 TCACGGGTTT AATCTTTTGT GTTTAGGGNN NNCACGGTTC AGACCAAATG GTATATAATT 360
 TTN 363

SEQ ID NO:7640

SEQUENCE LENGTH:270

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08887

SEQUENCE DESCRIPTION:

GATCTGGTTC TGAGGAGGAC ACACCTGGCA TCGGATGACC TTTATAAATT ATCTATGAAA 60
 ATGCCAAAAG CTCTCAAGCC AAAGAAGAAG AAAAATATTT CCCATGATAC TTTTGGTACA 120
 ACTTATGGAA GGATTCATAT GCAGAAGCAA GACCTAAGCA AACTACAAAC CAGGAAAATG 180

EP 0 679 716 A1

AAGGGGTTGA AGAAGCGACC TGCAGAAAGG ATAACAGANG ACCACGNGGA AAAAGTCAAA 240
AAGAATTAAG AAAAATTGAT GGAACCTAAA 270

5 SEQ ID NO:7641
SEQUENCE LENGTH:336
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS08888
SEQUENCE DESCRIPTION:
GATCTCTGAA AAGACGTTAT CACCTTAAAG CTCAAATTCT TTGGGATGGT TTAACTTAA 60
GTCCATTAAAC AATTCAGGTT TCTAACGAGA CCCATCCTAA AATTCTGTTT CTAGATTTTA 120
AATGTCAAAGT TCCCAAGTTC CCCCTGCTGG TTCTAATATT AACAGAACTG CAGTCTTCTG 180
15 CTAGCCAATA GCATTTACCT GATGGCAGCT AGTTATGCAA GCTTCAGGAG AATTTGAACA 240
ATAACAAGAA TAGGGTAAGC TGGGNTAGAA AGGCCACCTC TTCACTCTCT ATAGANTATA 300
GTAACCTTTA TGAACGCGGG CCNTNTAGTT TGGGTN 336

20 SEQ ID NO:7642
SEQUENCE LENGTH:331
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08889
25 SEQUENCE DESCRIPTION:
GATCCATTTT AGATACTTGT GGCAGCAAAC AAAGCAGTTC ACCTCTACAA ACTGGGAAAA 60
ATGAAGACAA GAACTCTATC TACTGAAATN ATTTTCANCC TTCCCCCAA TAACAATATT 120
TCAGAGGCTT TGAAAAAATT TGGTATCTCA GCAAATGACA CTTCAATTCT AATTGTTTAC 180
ATTGAAGAGG GAGAAAAACA AATAANTCAA GANTACCTAA TATCTCAAGT AGAAGGTCAT 240
30 CAGGTTTCTC TGAAAANTCT TCCTGAAATA ATGATTATTA CAGANGTCAA ANAGGTATAT 300
AANCTCTCTT CACAGGNGGN ANGTATCGGA N 331

35 SEQ ID NO:7643
SEQUENCE LENGTH:327
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08890
40 SEQUENCE DESCRIPTION:
GATCCACTGT CCGGCAGAGG AAAAATGCCA GCTGTGGGAC AAGGAGTGCA GGCCGCACAA 60
CCTCGGCAGG CACCGGGGGG ATGTGGCGAT TCTACACAGA AGATTCACCT GGGCTCAAAG 120
TTGGCCCTGT TCCAGTATTG GTTATGAGTC TTCTGTTTCT CGCTTCTGTA TTTATGTTGC 180
ACATTTGGGG CAAGTACACT CGTTCGTAGA TTCAGTTACA TCCATCTGTC ATCTGAAGAA 240
45 GGAGGAAAAA ACCCAACATT TCTTGGACCA AAAGTATAGT GACTATCTGT TCATGAGAGA 300
AATTTTCTGT NAGCTTGCTG TTTTGAN 327

50 SEQ ID NO:7644
SEQUENCE LENGTH:287
SEQUENCE TYPE:nucleic acid

55

TOPOLOGY:linear

CLONE:HUMGS08891

SEQUENCE DESCRIPTION:

5 GATCAAGATG AAGTTCAGCT AGAAGTCATA CCACCCTCAG GAATCAGCTA AGTAATTATN 60
 ACTTGATTCT TTTAGCAAAT CAATGCACGT NATCCTACTT AATCCTTAAA TAAGTTTAGA 120
 TTAACTAAC CCAAAGTCCA GGAGGATGTT CTTACAAAAN TAGCTATATC ANGGGCTGGC 180
 ACCTAGACAT TAANCTGTAA TTNGANAATA AGCAACATGT TGCATAACTN GTTGAATAA 240
 10 TTCCTTGTCN TGTTTAACAC TTGCATAAN TTAGCAGNGT AANNNTN 287

SEQ ID NO:7645

SEQUENCE LENGTH:327

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS08892

SEQUENCE DESCRIPTION:

20 GATCTACCCA ATTAGAAATG CAACTTGAGA AACCCAAACC TGTAACCA GTGACGTTTT 60
 CCACAGGCAT CAAATGGGT CAACATATTT NACTGGCACC TATCACAAG CTTGAAGAAG 120
 CTCTGTATGA ATACCAGCCA CTGCAGATAG AGACATATGG ACCACATGTN CCTGAGCTTG 180
 AGATGCTAGG AAGACTGGG TATTTAANCC ATGTCAGAGC TGCCTNACCA CAGGACCTTG 240
 CTGGAGGCTA TACTTCTNCT CTTGCTTGTC ACAGNGCACT ACAGGATGCA TTCNGTGGGC 300
 25 TTTTCTGGCA GCCCAGTTAA CCATTTN 327

SEQ ID NO:7646

SEQUENCE LENGTH:329

SEQUENCE TYPE:nucleic acid

30 TOPOLOGY:linear

CLONE:HUMGS08893

SEQUENCE DESCRIPTION:

35 GATCTGGCTG CTCATGAAGA CAAAGTTCTG AGTGTAGACT GGACAGACAC AGGGCTACTT 60
 CTGAGTGGAG GAGCAGACAA TAAATTGTAT TCCTACAGAT ATTCACCTAC CACTTCCCAT 120
 GTTGGGGCAT GAAAGTNAAC AATAATTGA CTATAGAGAT TATTTCTGTA AATGAAATTG 180
 GTAGAGAAC ATGAAATTAC ATAGATGCAG ATGCAGAAAG CAGCCTTTTG AAGTTTATAT 240
 AATGTTTTCA CCCTTCATAA CAGCTAACGT ATCACTGGGG CTTATTTTGT ATTTTATAAT 300
 40 AAGNTAGGGT TGTGTTTTNT NAAATNCGN 329

SEQ ID NO:7647

SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

45 TOPOLOGY:linear

CLONE:HUMGS08894

SEQUENCE DESCRIPTION:

50 GATCTTCAAG TGAACATCTC TTGCCATCAC CTAGCTGCCT GCACCTGCCC TTCAGGGAGA 60
 TGGGGGTCAT TAAAGGAAAT CTGGGACAAC AAA 93

55 SEQ ID NO:7648

SEQUENCE LENGTH:283

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08895

SEQUENCE DESCRIPTION:

GATCTGCATG CATTGCTTGC ATTTTNTTGG TATCTGAATG TTGGTTCCTT GTTCCAGGAA 60
 TTCAACATTA NTTTCCAAAA GTATCATGGG ACTTGTGACA ATACANGTCA TGAATCTATG 120
 TATAAAATTT ATCGGCCTTT CTCATTACC TGCTCTAGTA TTATTGTATT GTGTGTGCGT 180
 GCGTGTGTGA TGTCAGGCTG CCACGTAAAA CTTAGAGGA AAATCTTAA NGCAGGCCAT 240
 CCTTTTNGC ATGCTCTATT CTAAGTAGGA ATGTTCAATG TAN 283

SEQ ID NO:7649

SEQUENCE LENGTH:276

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08896

SEQUENCE DESCRIPTION:

GATCACCTGT AGGTCAATAG TTCGAGACCA GCCTGGCCAA CAAGGTGAAN CCCATCTNTA 60
 CTAAAAATAC AAAAAATTAG CCAGGTGTGG TGGTGTGCAC CTGTAGTCCC AGCTATTTGG 120
 GAGGCTGAGG TGGGAGAATC ACTTGAACCT GGGAGGCGGA GGTTCAGTG AGCTGAAATC 180
 GCGTACCACA CTCCAGCCTG GGTGACAGAG CGAGACTGTG TTTCAAAGAA AAAAAAACC 240
 AGGCAGCCTT TTGCTTGGTT GGAATCTGAT TTTCTN 276

SEQ ID NO:7650

SEQUENCE LENGTH:182

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08897

SEQUENCE DESCRIPTION:

GATCTAAATA TTCAGGTTTT AAGCCTGCTG CAACTTTTA AAATATTATG ATAGNTTCTG 60
 TACTACATGT GGGAAACAAG CAAGAACTAA ATAATCAAAT GTTGTCAACC AAAAGTAATA 120
 GTTGGGTATT GGAGATTTTT TAAAATGTT TTTATGTTAT TNGCTATTTG GAGTTAAATA 180
 AA 182

SEQ ID NO:7651

SEQUENCE LENGTH:93

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08898

SEQUENCE DESCRIPTION:

GATCTNAGAA AACGTTTAAT ATCAGATTN TTTATTTTIN CCAACACAAC TGAATCAGTT 60
 TTCTTAAATA AAAGTTTATT GTCTACTGTT AAA 93

SEQ ID NO:7652

SEQUENCE LENGTH:275

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08899

SEQUENCE DESCRIPTION:

GATCTAAAGT TAGTTGCCTT TGCCTGTAAA ACATGTGATT TGCAAATTAT TATTTNNNTT 60
 TTTTAAACA AATGGAAGTA AATTGTTTC ACGTAAATCT TAATTTTCAA CCTTCTGGA 120
 TACCTTAATT GTAACGTCA GTTGCACGTG GTCGGTATAT GGAAACACAT TGCTCTACCC 180
 TGCTACTTAG TTGNTTTTAA AGTGAATTN CAGTGATGNG AAATTGTGA AANNTATATT 240
 GTATTTCTTT TGATGTTTCA AAAGGTTGCC TATGN 275

SEQ ID NO:7653

SEQUENCE LENGTH:279

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08900

SEQUENCE DESCRIPTION:

GATCTTTAGG TGTGAACTG GGAATATGCT CTGCAACACC CAGGATTAGG ATAGAGATGG 60
 ATTCATTTT ATATGGGCTA GCATATTATT ATTAAAGACC ATTCAGGAG TGGGACTGTG 120
 AAAGGAGGTG CTCTGTTATC TGGTCCAAGA GCTCCTCTGG GAATTAATCC TGGCTGGTTC 180
 TGATGAGCTG GCAAGGNGCA GNGGATTGCA GTCTTTAAGT AGAAAATTNG AACCCCATCC 240
 ATATAACACG TGNACGTGCA TTAAATAGCT TGAATTGTN 279

SEQ ID NO:7654

SEQUENCE LENGTH:264

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08901

SEQUENCE DESCRIPTION:

GATCAAGACC AGAAAGAAAT TATTGATACC AATGGAGCTG GAGATGCATT TNTTGGAGGT 60
 TTTCTGTCTC AACTGGTCTC TGACAAGCCT CTGNCTGAAT GTNTCCGTGC TGGCCACTAT 120
 GCAGCAAGCA TCATAATTAG ACGGACTGGC TGCACCTTTC CTGAGAAGCC AGACTTCCAC 180
 TGATGGAAGA GCTGAAAACA CAAGCCCAGG AGTGCAGACA CTGCCCTAAT TGCTTCCTGA 240
 GAATTCCTCAT ATTAATAAAG ANGN 264

SEQ ID NO:7655

SEQUENCE LENGTH:61

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08902

SEQUENCE DESCRIPTION:

GATCTTTTAG GAAAAAAAT AAATATTTT AACATTAAAG CAGTCACTTT GGTATATCAA 60
 A 61

SEQ ID NO:7656

SEQUENCE LENGTH:66

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08903

SEQUENCE DESCRIPTION:

GATCTGTTTG CTTAGCTGTC AACAAAAAGA AACCTGAAG GAGCATTGG AAGTCAATTT 60
GAGGTN 66

SEQ ID NO:7657

SEQUENCE LENGTH:252

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08904

SEQUENCE DESCRIPTION:

GATCTNACTA AGCTTTCTTA GGAAGGGAAA GCATCTGGCA AGGAGTAAAA TGTCTGTTTC 60
CAACAATTGT CCCTGTTCCC ACTACTTTTT AAAAATGTCC TTAGTAAAAA GTGTATATGA 120
GGCAAGATTA TATTGAAAAA NTGAATATAA GAAAAAATA TTATGTTAGA GAGNAACGTA 180
TTCAGCATTG TTNTCAAAAT CACAACTGT TAGGNGACTC TGACATCAGG TTTGTTCTCC 240
ATATGCAGCA AN 252

SEQ ID NO:7658

SEQUENCE LENGTH:153

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08905

SEQUENCE DESCRIPTION:

GATCTACATG GATGTTCTGT TCTCTGAACT GTCTGGATGA ACCGGTCAAC GGCACATC 60
ATACCTTAGT TTTTAAATCT GCATTGTGGT CATAATCTGT TATTTAATTA ATTTCTCGTA 120
TTTTTAATAA AAACTTTGCC TATATATTTT AAA 153

SEQ ID NO:7659

SEQUENCE LENGTH:23

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08906

SEQUENCE DESCRIPTION:

GATCCCAAAA GTAGCAAGAG AAA 23

SEQ ID NO:7660

SEQUENCE LENGTH:280

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08908

SEQUENCE DESCRIPTION:

GATCAATTGC AGAATGATAC AGATAGTACA ATGACTTTTT TGTAATTG AATTAAAAAGT 60
TTCCTCATAA AACTCAGTAC TATCTATTGG TAATGGATGC ATATATGTAT GTACATGTTT 120

TTNAAACGGA TTGGAAGGAT ACAGACCAAA CTCTTGATAG TGGTCACCTG TGAAGAGTGG 180
 AGAAGGGAAA ATNATGAGTG TGGGAGGTGG GATGGGTATT GGTAAANGGG GACTTCAGCT 240
 TTTTATATAA ACATCCACTT CTCTTTCAAA AAGNCTTCAN 280

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SEQ ID NO:7661
 SEQUENCE LENGTH:137
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS08909
 SEQUENCE DESCRIPTION:
 GATCATTTTA AAAAAGTGA TCTATATNAT NTTAACTTC AGAAGGCATC ATTTATAAGA 60
 CAGTATGGCA GTTAATTATA AAATNATTTT GATGAATTAT GATACAATCT ACATAATAAA 120
 GAATNCTTTT GATTAAA 137

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SEQ ID NO:7662
 SEQUENCE LENGTH:305
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS08910
 SEQUENCE DESCRIPTION:
 GATCCTAAAA CTGTCTCTCA CATTATATAG TAGATGTTTG TTTATAATGT TTACAAAACA 60
 TTTTGGTGAA TTTCTCAAT GTTTTATAAA TGTACATTTT TTAAGTCCTT AAGCTGACTC 120
 TTAGCCATCA TGTAGCTTAA GGAGTCTGAA ATCTGCCATT AAAACTGCAC CTTTAAGCCA 180
 GGTGTGGTAG CATGTGCCTA TAGTCCCAGC TACTTGGGAG GTGGAGGTGG GAGGATTATA 240
 AATAGAGACT TTCCTAAGA CTTTAAAAAT GTATTGNNA ACTATTTTTT ATTAAATACT 300
 TTGTN 305

30

SEQ ID NO:7663
 SEQUENCE LENGTH:103
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

35

CLONE:HUMGS08912
 SEQUENCE DESCRIPTION:
 GATCCTAACC TTGAAGTATG CCTTGAACCT ATTAACATGG CCATTATAAG AATAAAATAT 60
 GTAGTTGTGT CTTAATGGAA TTAATAAATG TCATTTCACCT AAA 103

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SEQ ID NO:7664
 SEQUENCE LENGTH:211
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

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CLONE:HUMGS08913
 SEQUENCE DESCRIPTION:
 GATCAAGGAA TAAAGAATCA TTAAAGAATG GGGTGTGGG GGTTCCTTA ATTATCAGTG 60
 ACACATTAAT TTTTTCTT AGAGCTTCAT ACAATAACA CATTGCAGAA AGTTTtagAA 120
 AGTTTTGCAG GACCTTAAAT TTTCTGCCCT GACACTTTTC GTCATGTTTC CTGTATTCCN 180

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GGTTATTGTA GCGCATGCTT TTCAGGTAGT N

211

SEQ ID NO:7665

SEQUENCE LENGTH:71

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08914

SEQUENCE DESCRIPTION:

GATCTTAAAT TTTGTGATAT GGAGCCCTGT AATNTTTTTC TTATATAAAA ATGGGTATCT 60
 ATATTCATAA A 71

SEQ ID NO:7666

SEQUENCE LENGTH:344

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08915

SEQUENCE DESCRIPTION:

GATCCCAGCT CATGGCAGAG CCGTTTGAT CCTCGGAAGA ACGGAAGGAA CGATGGGAAC 60
 AGGGCCAGGC TGATTATATG GGAGCAGATT CCTTGACAA CATCAAGAGG AAACCTTGACA 120
 CTTACCTCCA GTAGAAACAC TGCATTTTTC TGTGAACACA TCCACTTCAC AAGCCTTGTT 180
 TCTGATACTT AGTATCTAGA GCTGGGTTGA GAAAAGTCTG TTACAGTTGC TAGAGGTTTT 240
 CATTAAANCT TATCAGGNGG GCGGCTTTTT TAGGATAAGA GGTGAGAACT GGGCAAAAGT 300
 TGTGAAGCAG CAATTCTGTT ATATGGNCAG TGTCTGCTT TTTN 344

SEQ ID NO:7667

SEQUENCE LENGTH:200

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08916

SEQUENCE DESCRIPTION:

GATCAGGACT GCTGCACTCC AGCCTGGGTG ACAGNGCGAG ACTCCATCTA AAAAAACAAA 60
 AAGCAAAAAA ACCCACAAAT ACCTCATGGA GATGAACTGT AATAATTGCT TAAAGTTCCA 120
 TTAAATTATG TTAACCTAA TCTAGCAAAA ACATAGATGT ACTTAAAAAT AAATCATGGA 180
 TAATGATTTT TTAACCTAAA 200

SEQ ID NO:7668

SEQUENCE LENGTH:277

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08917

SEQUENCE DESCRIPTION:

GATCTCATTT TCAAGTNTGA GTCTGACCAA CCATGGAAAA TATTCGACAT GAATTAATGT 60
 AGAGAACTAT AAAGCATTTA TGACAGCTCC AAGAAAAGTC ATCTACTCTA TGCAGGAGAT 120
 ATGTTTAGAG ACCTCTCAGA AAAACTTGCC TGGTTTGAGG GTACACAGTA CCATTTTAAT 180
 CTTCTGAAAA TATCTGTATT CCTGCTCTGG GGNTGCTGTC ACTGTCAATC TGCTATATT 240

TTCACATATCC TATTAAAATA TTACTGTCTC CTTTAAA

277

SEQ ID NO:7669

SEQUENCE LENGTH:163

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08919

SEQUENCE DESCRIPTION:

GATCTTCTAC ATTTATATTT TTAATCTTCT GTTAAATACA CTTTCCGATA TTGCCTTGCC 60
TTTGAGCTC TTGCTACAGT CGCCTTTGCT ACTGCTTTAA GAGAATTTAC AGGTATTGAT 120
AAAGAACAAG ACTGTTTTAT TAAAAGNTTT ATTCAACTTG AAA 163

SEQ ID NO:7670

SEQUENCE LENGTH:111

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08920

SEQUENCE DESCRIPTION:

GATCCAAATC TGGTTCAAAC ATTCAAACT TCAAAGATAA TTCATCTTTC GCTAATGCTT 60
GTGTTTCTGT NGTTCCCTTG AAAAAAATA AAAACCCGAC CCAGCAGCAA A 111

SEQ ID NO:7671

SEQUENCE LENGTH:155

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08921

SEQUENCE DESCRIPTION:

GATCTGTAC TGTTCCTCGT CACCCTCAGA TGGGACCGTC TAGTTGTAGG AAAACAAGCT 60
CAGGGCTCCC ACTGATTCTA CATTTTGGTG AGTTGTATAA TTGTTATGTA TTGCAATGTA 120
ATAATAATAG AAATAAAGTG TACGATAAAT GCAAA 155

SEQ ID NO:7672

SEQUENCE LENGTH:239

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08922

SEQUENCE DESCRIPTION:

GATCTGGAGA GGTGTTTGT GAAAGTGGAG AAGATTCACG AAAAGGTCTA AATGAATTGC 60
GTGTGCAGGG CGCGGATTTA AAGTCCAATT TCTCATGACC AAAAAATGTG TGGTTTTTTC 120
CCATGTGTCC CCTACCCCCC AATTTCTTGT CCCCTCTTAA AGAGCAGTTG TCACCACCTG 180
AACACCAAGG CATTGTATTT TCATGCCAG TTAACCTATT TACAATATTT AAGTCAAA 239

SEQ ID NO:7673

SEQUENCE LENGTH:327

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08923

SEQUENCE DESCRIPTION:

5 GATCCTTTCA TTAATCAGAA CAGCAGCTTA TNAACTGCTT CATTTTCCAA AAGTGTTTCC 60
 CAAAAATCAG AAACCTAAAC ACCCAACAAC AGGAATTATT GCCATCACAT TGGCGTTTTA 120
 CATATGTCAC GAAGTTCACC TAGCTGGTTT TAAATACAAC TTTTCTGANC TCAAGAGTCC 180
 TTTGCACTAC TATGGGAATG CCACCATGTC TTTGATGANT AAGAACGCGT ATCACAATGT 240
 10 GACTGCAGAG CAGCTCTTTT TGAAGGACAT TATAGAAAAA ANACCTCGTA ATCAACTTGA 300
 CTCANGATTG ACTCTACAGA CTCGGAN 327

SEQ ID NO:7674

SEQUENCE LENGTH:377

15 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08924

SEQUENCE DESCRIPTION:

20 GATCTCCCNT AATTGGGAAT GAANGATTTC ACAGACTAGA GTCTCCGATG CTGGTCATGA 60
 TGTCAAAAC AAGTTCTGAC TCATTTAGGG AACTGGATAC TTGGGTCTCC AGAAGGGCCA 120
 ATGGGAGGGC CATAATTCTG TTTATTTTCA AATTGTCTTG TTTTACCTT GTTAGAATGA 180
 NCTNTGGAAG CCCAGCCAGG GACAATGCAC CTTACAGAG ATTCTGCACT AATCTGAGTG 240
 AAGGTCTAAG GTTTGGAATC TCCCCTCAT GGAGAGAAGC TTTGTATGGC TGTCATGCTT 300
 25 AGACAGTGAT TCCTGCAACT TGACCTTCAG GCTGGGAGAG GTNGAGAGCC ATGCCTGTTT 360
 CTCCTTNCTT TGCTATN 377

SEQ ID NO:7675

SEQUENCE LENGTH:350

30 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08925

SEQUENCE DESCRIPTION:

35 GATCTGCTTC CCGCAAGACT CACAGCTGTT GGCAGGAGAC CTCAGTTTGT TGCCACATGT 60
 TCCCCTCCAG AGGGCTCTC ACAACATGGC AGTTATTTGT CCCCAGAGCA AGCAACACCG 120
 GAGGGCAAGG AAGAAGCCAT GATGTTTTT GTAACCTAGC CTCTGAAAGT GTCATACCAA 180
 TTCTGTATTT TGTTGGTCAC ACAGACCANG TCAACTACAA CGTGGGAGAC TCCTACACAA 240
 GGCATGAATT CTAGGAGGTG GGCATTTTTA AGTGTCTCT GGAAGGAGGC TGTCACAACC 300
 40 TGGAAGTTAA AAGCATTGAT ATTCTGAAAT ACAGCGTGTA TTACCATGGN 350

SEQ ID NO:7676

SEQUENCE LENGTH:335

45 SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08926

SEQUENCE DESCRIPTION:

50 GATCTAGAGG AGGATTTGCA GGAAGAGCTC GTGGAAGAGG TGGTGGCCCC AGTCAAAACT 60
 GGAACCAGGG ATATAGTAAC TATTGGAATC AAGGCTATGG CAACTATGGA TATAACAGCC 120

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5 AAGGTTACGG TGGTTATGGA GGATATGACT ACACTGGTTA CAACAACTAC TATGGATATG 180
 GTGATTATAG CAACCAGCAG AGTGGTTATG GGAAGGTATC CAGGCGAGGT GGTCATCAAA 240
 ATAGCTACAA ACCATACTAA NTTATTCCAT TTGCAACTTA TCCCAACAG GTGGTGAAGC 300
 AGTATTTTCC AATTGAAGA TTCATTTGAG GGNAN 335

10 SEQ ID NO:7677
 SEQUENCE LENGTH:295
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08927
 SEQUENCE DESCRIPTION:
 15 GATCCAAATT TGAAGCTCCA AGGATAGGAA AAAGCTTACT TGTGCCTGTC CTAGACAAGT 60
 ACTAGAGTTT ATATTGAAAT: CAAAACCTCT AAAAATAAAA ATCAGTAAGA TTAATTTTAC 120
 TCATTTTCT AACACGAGTC AGTATGCTTG ANGTTTTCAT TGTATTTGAT GTCATGTAAA 180
 TTTACTGTAA AAAATAAGAA AANNTTTTAA AATGCTTTAT TGATTAANTT TAGGGTTGCT 240
 TTTCAGTATA CTCAGTAACT CCGAAGTCAG TGTTAACAGT ACTGAGCTTA CATAN 295

20 SEQ ID NO:7678
 SEQUENCE LENGTH:136
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08928
 SEQUENCE DESCRIPTION:
 GATCTTTTGT TAACTTTTAA ATGGCCTGAG TCCATAGAAA CTGCATCCTG AGAGGTTGCT 60
 CGGCTAGAAA GCGGAAAACC GCTTCACATG AGTGCTCTAC TTTGTAATGA AAATAAACAC 120
 30 TATTAATACA GGCAAA 136

35 SEQ ID NO:7679
 SEQUENCE LENGTH:310
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08929
 SEQUENCE DESCRIPTION:
 40 GATCTGGGAA CGGAAGATGA ATGAATGGAA GATGAATGGT TTCTCTCCCA GTTTCTCTG 60
 AGGCAAACAT AATTATACCT NCTCCTTTGT TCTGGGAGGT TTTTCTCTC TCATCTCCCC 120
 TCCATTACAC AAAACAGAGT TGGGTCTATA TTTNACCCGG CTGGCTCTTG GAACTGCAGG 180
 GAATCCTGGC TGTTTTTATT TTTAAAAAAT TATGAGATTA NCTACCAAAC TTGAGATGTA 240
 AGTTGTGGGT TAGCTCTCGG ATTCTGCTAG ATAACCTGGA ACAAAGCATT AACTTCTTTG 300
 45 AGAGCTTCN 310

50 SEQ ID NO:7680
 SEQUENCE LENGTH:177
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08930

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SEQUENCE DESCRIPTION:

GATCTGCCAT GTAGGAATGA GACAAATACA GTTTGCTTAT GAAAGGAAAG TGGCATACTT 60
 TTAAATTTGGT CTACACAGAA AAGTAAAAGT AAACATTTCA TTAAATAAG ATTCATTATC 120
 TAATAAATAT TGAGGGAATA TTTTCCTAA ATAAAAATTT TTCTGACTGC TAACAAA 177

SEQ ID NO:7681

SEQUENCE LENGTH:84

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08931

SEQUENCE DESCRIPTION:

GATCTAAGCT TTCTTTAATA TAAAAAAAT NATACTTCTC TGAAGTGTTC AATAGAGCAA 60
 TAATTAAATA CATCTTCTGA TAAA 84

SEQ ID NO:7682

SEQUENCE LENGTH:285

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08932

SEQUENCE DESCRIPTION:

GATCTGTTGG GAAAGATAAG AAAGCTATTC AGGCATCAAT TAGACGTAAT AAGGAAACCA 60
 ACACCGTTTT GGCCAGATTG AATAGCGAAT TGCAGCAACA ATTAAGGAT GTTCTTGAGG 120
 AGAGAATTC CCTGGAAGTT CAACTGGAAC AACTTCGACC ATTCTNTCAC CTATAAGCCA 180
 ATTGCCGTTA ACTGTGAACA TACTTGTTTT TAAGTGGTTT TGGGTTCAAA GCCAATTTGG 240
 AGACCTAGAC ATTCAGCTCA CTGCTTAACT CACATTAAA TTTTN 285

SEQ ID NO:7683

SEQUENCE LENGTH:313

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08933

SEQUENCE DESCRIPTION:

GATCTCAATT CTGTTTCCTC TCACATGATT ACTTGATAGC TAAGCATCTG ATTGGTTTAC 60
 TGCTTTACCA CTGAGCTGAA ATGCCGTGTT TTCCATTTAT TAAATCACA CATGGCTCCT 120
 GTTTTNTCA CTCAGCACTT TTTCTCCATA TTCTTCAAGA CGATTGTGAG TATGGTACGT 180
 AACAGGAATT ACATCTGGTA AGTTGTATAG TTTTGTGTAG GAACTCTATA TTCATAGCAT 240
 ATTTGTGGAA ATGATACCTA TGGAGGTTTC TCACACTGGT GTGTCATTAT ACATTAATTG 300
 TACAATATGC ATN 313

SEQ ID NO:7684

SEQUENCE LENGTH:110

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08934

SEQUENCE DESCRIPTION:

GATCAGTAAT GGCAAGAGCC TTTNATTCTC GAATGTTTAA AGCCTAGGAG TTCTACAAAA 60
 TTGGNTTCTT TCTACAAGAN TCCCAAAATG GAATGCCTAA AGAGGTNTNN 110

5 SEQ ID NO:7685
 SEQUENCE LENGTH:132
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08935

10 SEQUENCE DESCRIPTION:
 GATCCCAAAA ACCGGGCAGC CCAGGAAGGA ACTGGGGAAG GTGGTCATTC AGGGGAAGAA 60
 CCAGGATGCA GGGCTGGCTC AGGGTCTGCG CAAGATGTTT GGCTGATTAA AAGTTAAACC 120
 TTAAGAGAGA AA 132

15 SEQ ID NO:7686
 SEQUENCE LENGTH:85
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08936

20 SEQUENCE DESCRIPTION:
 GATCCCATGC CTGAAATNTG GAAGCATATG TACAAAAATC ATTTTACGT TTTATTTTAA 60
 ATAAATCATT GTGTTTGACC GTAAA 85

25 SEQ ID NO:7687
 SEQUENCE LENGTH:333
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08937

30 SEQUENCE DESCRIPTION:
 GATCCTAGTC ATGACTATAC ATTGTGACAT GGTCATTACA TATGGATTAG ACCAACTGGA 60
 GAATTGCCAG ACTTGTGGTA CCGATTATAT CATCTCAGTC TTGAATTAC TCACGCTGAT 120
 35 TGTGAAACAG ATAAATACGA AACTGCCATC ATCATTTGTA GAAAAACTGT TTATACCATC 180
 ATCTAAACTA CTATTCTTGC GTTATCATAA AGAAAAAGAG GTTGTGCTG TAGCCCATGC 240
 TGTTTATCAA GCAATGCTCA GCTTGANGAA TATTCCTGTT TTGGAGACTG CCTATAAGTT 300
 AATATTGGGN GANATGACTT GTGCCCTAAA CAN 333

40 SEQ ID NO:7688
 SEQUENCE LENGTH:333
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08938

45 SEQUENCE DESCRIPTION:
 GATCTGATTC GCTCCATCAA TGACCCGGAG CATCCACTGA CGCTAGAGGA GTTGAACGTA 60
 GTAGAGCAGG TGCGGGTTCA GGTAGCGAC CCCGAGAGTA CAGTGGCTGT GGCTTTCACA 120
 CCAACCATTC CGCACTGCAG CATGGCCACC CTTATTGGTC TGTCCATCAA GGTCAAGCTT 180
 50 CTGCGCTCCC TTCCTCAGCG TTTCAAGATG GACGTGCACA TTA CTCCGGG GACCCATGCC 240

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EP 0 679 716 A1

TCAGAGCATG CAGTGAACAA GCAACTTGCA GATAAGGAGC GGGTGGCAGC TGCCCTGGAG 300
AACACCCACC TCTTGGAGGT TGTGAATCAG NTN 333

5 SEQ ID NO:7689
SEQUENCE LENGTH:110
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08939
10 SEQUENCE DESCRIPTION:
GATCCATGAA CCCTTTTAT AAGCTGTGTG TGTCTCTGT ATTATTGTA TTAAGTATT 60
TTAGNATTT GCCTGTAAGT TATTAAAGAC TGATAACTGT AGCTCTTAAA 110

15 SEQ ID NO:7690
SEQUENCE LENGTH:186
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08940
20 SEQUENCE DESCRIPTION:
GATCCAGCCA AACGAATTC TGGCAAAATG GCACTGAATC ATCCATATTT TAATGATTG 60
GACAATCAGA TTAAGAAGAT GTAGCTTTCT GACAAAAAGT TTCCATATGT NATGTCAACA 120
GATAGTTGTN TTTNNTTGT TAACTCTGT CTATNTCTCT CTTATATATA TCCCNCTNTT 180
25 ATCAAA 186

SEQ ID NO:7691
SEQUENCE LENGTH:231
SEQUENCE TYPE:nucleic acid
30 TOPOLOGY:linear
CLONE:HUMGS08941
SEQUENCE DESCRIPTION:
GATCATGTGA CCAATTGTGC ATTCAGTTTT CAGAATTCTT TGCTATATGA TTTGGATTAA 60
TTCTATATAA TTTTGGACTT TTAATATTA AGGTAAAAA ATACCTGTAT CTAAGATTGA 120
35 TTCTGTTAAC TGTGTCTTA AACTAAAGG TATTAAAGTA TAANNTTAAA ATTTGCAATT 180
TTTTTAAAN NNTTGCAATT TTGATTCTCA TGGGGGAAAT TGGAGATAAT N 231

40 SEQ ID NO:7692
SEQUENCE LENGTH:277
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08942
45 SEQUENCE DESCRIPTION:
GATCACAATT TAATAACAAG TAAACTTTA CTTTGACAA TTTTGCCACT TATTTTGTA 60
TTTGTAAACA CACGTTTGA GATATATTA TGCCATTAAT ATTTAAGGC ATGTTTAAA 120
ATTTACTGTT TTCTGAAAAA TTAATGNNN CTGGTTTGAT TGCTGGAGAT TATTCCAGT 180
TTTNTCTAAA TGTATTTNA TNCTGTATGG TACATTTNG NGGATTNCCT GATTAATGTC 240
50 ATCTTAGCAT TATATTCTGG TTATCTATAA TNNANN 277

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SEQ ID NO:7693

SEQUENCE LENGTH:312

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08943

SEQUENCE DESCRIPTION:

GATCTGGGCA AAGCAAATTG AAAACCTTCT GGAAAGGATT CACCATTTTA GATGCCATTA 60
 AAAACATTCTG TGATTCATGG GAGGAGGTCA AATTGTCAAC ATTAACAGGA GTTTGGAAGA 120
 AGTTGATTCC CNNCCTCATT GATGACTATN AGGGGTTCAG GACTTCAGTG GAGGAAGTAA 180
 GTGCAGATGT GGTGGAAATA GCAAAAGAAC TAGAATTAGA AGTAGAGCCT GAAGATGTAA 240
 CTGANTTGCT GCAATCTCAT GNTAAANCTT TAACAGNTGA GGAGTTGTTT CTTATGGNTG 300
 CGCAAAGAAA TN 312

SEQ ID NO:7694

SEQUENCE LENGTH:83

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08944

SEQUENCE DESCRIPTION:

GATCTTTACT GTTCAGAATT TAGGAAAGTT CTCTGGCTGT TGCATCCAAG TAAAATTAAA 60
 ATAAAATTGG TTGCANNITT AAA 83

SEQ ID NO:7695

SEQUENCE LENGTH:263

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08945

SEQUENCE DESCRIPTION:

GATCCCTTCC TCTGTCTCTC TAACTCCCCA GCCATGAATT TTGGCCCCAG GGAGAGGAAT 60
 AGGCAGTATA GGAGGGATGT CAGGGAATC ACATCTGTGG TTGAATCTCA GAAACAATGG 120
 AAAGGCAGTT GTCTTTGGAT ATGGTGAAAT TGTGGTAGAG CACCCATGGT GGGGGTCTGG 180
 CAGGAGCTCT GCGGCTTCGG TAGCAGCAGT ATGAGGAGAG CTGGGCCACA TGCTTATGGG 240
 TTAGGAGAAG GTAATTTCCA GTN 263

SEQ ID NO:7696

SEQUENCE LENGTH:142

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08946

SEQUENCE DESCRIPTION:

GATCATGGCT GAAAAAGCAG CAAATGCAGC AGGAAAAAG TTCCGAAAGA AGAAGAAATT 60
 TCGCAATTAA GATTACCAA GCAAAGTCA ACATTTTACA TTGCTCCTT ATTTACTTAT 120
 TAAAGACGTT TGGAAACTA AA 142

SEQ ID NO:7697

SEQUENCE LENGTH:200

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08947

SEQUENCE DESCRIPTION:

GATCAAGATT GAGAAGACTA GATACATAAC AACCAACTGC AATGTTTGGT CCTTCATTGA 60
ATGGTTGAAC AAACCAGCTG TGAAAGACAT TTTGGGACAT ACAGACATGG ACTGAGAATT 120
AGACATAAGA AAATTATTTT GTTAACTGTT ATCATGGTTA ACAAACGGT CATATAATNT 180
GATTATGTAG AGAAGTAAA 200

SEQ ID NO:7698

SEQUENCE LENGTH:351

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08948

SEQUENCE DESCRIPTION:

GATCTGCCTG CCTCAGCCTC CCAAAGTGCT GGGNTTACAC GCGTGANCAC CACACCCAGC 60
CTCCATCTTT AAACTTTAA ATGTGAAATT TCTATCATGT ACCGTTAGCC TAACAAGATT 120
TTNTTTCCTA TTTCTGACTG GTGCCTTTCC CCTTTTtagg AGCAACGAAA GCTACTCTCT 180
TAGTTATGTT CTTGTGATGT GACAAAATGT CAAGAAGATA GGAGAAGAGA ATATTTTATT 240
TCGTTGATGC TTTTGTTCCT AAGTGTGACC CTAACCTTAA GCTTTGTAGG AGTTGACATT 300
CTTTCATGTC CCTTCCCTTT ACTCATGCCG AAACCTATCAA CTGGGNCATG N 351

SEQ ID NO:7699

SEQUENCE LENGTH:350

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08949

SEQUENCE DESCRIPTION:

GATCTGTTTC TGGGGCCTCT GCGGGTGGCG GCCTGGGGCG GCGCGACGGC TGGTGCGCAG 60
GTACANTGAT GCTGAAGTAC TATGAGCCTT CGGAACTTGT GGAGAGACTA CAANGTTTGT 120
GNNNTATGG TCCCTTTAGT TGGGCTCATA CATTTGGGGT GGTACAGAAT CAAAAGCAGC 180
CCTGTTTTC AAATACCTAA AAACGACGAC ATTCCTGAGC AAGATAGTCT GGGACTTTCA 240
AATCTTCAGA AGAGCCAAAT CCAGGGGAAG TAGCAGGCTT GCAATCTTCA GGTAAGAAG 300
CAGCTTTGAA TCTGAGCTTC ATATCGAAAG AAGAGATTGA AAAATACCNN 350

SEQ ID NO:7700

SEQUENCE LENGTH:351

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08950

SEQUENCE DESCRIPTION:

GATCCCGCCA GAAATGAATA GACACCGCAA GAAACAAAGC TCGAGTGAAA GCTTGTTACA 60
TAATGATTGG ACTCACAATT ATCGCCTGCT TTGCTGTGAT AGTGTCAGCC AAAAGGGCTG 120

EP 0 679 716 A1

5 TAGAACGACA TGAATCCTTA ACAAGTTGGN NCTTGGCAAA GAAAGCTAAG NGGCGTGAAG 180
AAGCTGCATT GGCTGCACAG GCTAAAGCTA AATGNTATTC TAAAGTGACAA AGTGTTCACC 240
TGAATACCAT CCCTGTCATC AGCAACAGTN GAAGATGGGA AAAATNGAAT ATTTACCAAA 300
ATNTCTGCCA TGGTTTTATT TTGGTAACCN GATGCACAAT GTCTTTTTTT N 351

SEQ ID NO:7701
SEQUENCE LENGTH:85
SEQUENCE TYPE:nucleic acid
10 TOPOLOGY:linear
CLONE:HUMGS08951
SEQUENCE DESCRIPTION:
GATCATATGT GAAGATGTCA ATTAAGCTTG CATTAAGCCA CCTGCTTTGT AAGTGGATTG 60
15 ATTAATAAAT AACTTATATT TCAAA 85

SEQ ID NO:7702
SEQUENCE LENGTH:89
SEQUENCE TYPE:nucleic acid
20 TOPOLOGY:linear
CLONE:HUMGS08952
SEQUENCE DESCRIPTION:
GATCCCAGGG GCTTATCTCT TCAAGTGTGG AGAGGGCAGG GTCCACGCCT CTGCTGTAGC 60
25 TTATGAAATN AACTAATTGA AAATTCAAA 89

SEQ ID NO:7703
SEQUENCE LENGTH:375
SEQUENCE TYPE:nucleic acid
30 TOPOLOGY:linear
CLONE:HUMGS08953
SEQUENCE DESCRIPTION:
GATCCAGTGG CAATTGTAAA ACTAGCTATT TATGGCATGC TGCCAAAAAA CCTTCACAGA 60
35 AGAACATTN ATGGAAAGGT TGCATCTTTT TCCAGATGAG TATATTCCNN CNGATATTCT 120
TAAGAATTTA GTAGAGGAGC TTCCTCAACC ACGAAAAATA CCTAAACGTC TAGATGAGTA 180
CACACANGAA GAANTAGACG CCTTCCCAAG ATTTGTGGAC TCCACCTGGA AGATTATCGG 240
CTATAAGAGA ATAAGATTG CAGATAATTA ACAGTTGANG TGATTGAGAC TTTCTNCTTG 300
40 ATGAGTTTCT CTAACCTNCA GGGTGGTGT AACCAACTGC TACAGTTCAG CACCTGTTTT 360
TNTTGTGCCG NTTN 375

SEQ ID NO:7704
SEQUENCE LENGTH:276
SEQUENCE TYPE:nucleic acid
45 TOPOLOGY:linear
CLONE:HUMGS08954
SEQUENCE DESCRIPTION:
GATCACATCC TACTTCCTCA ATGAAGGGTC CCAAGCCCGT CCCCCTTCTT CCCACCGATA 60
50 TTTCTNGAA CGCGGCCTGG AGTCAGCAAC CAGCCTCTAG CAGCTGCCTC TACGCGCTCT 120

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EP 0 679 716 A1

ACCTGCTTCC CCAACCCAGA CATTAAAAATT GTTTAAGGAG AACCACACGT AGGGGATGTA 180
CTTTTGGGAC AGAAGCAAGG TGGGAGTGTG CTCTGCAGCC GNGTCCAGCT ACTTCCTTTT 240
GGAACCTTTA AATAGAATGG GTGTTGGTTG ATTAAA 276

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SEQ ID NO:7705
SEQUENCE LENGTH:220
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08955

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SEQUENCE DESCRIPTION:
GATCCACTGC CATTCTAATT GCTTTAACAA GTCATTACCA CACTACTGTT ACATCTTAAT 60
TATGCATACA GACAGGTAGA CTTGTTTTAC ATATGTGAAC TAACTAGTTG TCAAAGCAAA 120
TGCAGATTGT ATTCTGCAAG TAAAGTCTTT TTCTCTCTGA AATTCTAGG GATGTTCTTT 180
AAGTGAAATT CATATTAATA CTGAAGATTT TNGTTACAAA 220

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SEQ ID NO:7706
SEQUENCE LENGTH:252
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08956

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SEQUENCE DESCRIPTION:
GATCCAAAGT AACTNAAACA GGAGCCCACA AGACGGTCTG CCAGATTGTN AGCGAAACCT 60
GCTCCACCAA AACCTGAACC CAAACCAAGA AAAACATCTG CTAAGAAAGA ACCTGGAGCA 120
AAGATTAGCA GAGGTGCTAA AGGGAAGAAG GAGGAAAAGC AGGAAGCTGG AAAGGAAGGT 180
ACTGCACCAT CTGNAAATGG TGAACATAA GCTGAAGAGG CACAGAAAAC TTGNTTCTNG 240
TNGATTAACG AN 252

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SEQ ID NO:7707
SEQUENCE LENGTH:166
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08957

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SEQUENCE DESCRIPTION:
GATCTCCTGA CCTCAGGTGA TTCACCCACC TCAGCTTCCC AAAGTGCCGG GNTTACAAGC 60
TTGAGCTACC GCGCCAGCC AACACATCAC TTTTATCTTG CAGCTCCCGA GATTCTCTTT 120
TCTGTGACTN TAAAATTGT GCTNACATAT GTGTTTGATA TAAATN 166

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SEQ ID NO:7708
SEQUENCE LENGTH:52
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08958
SEQUENCE DESCRIPTION:
GATCAAAATA TTCTGGAAAA AAATAATAAA AAATAATTTG TATACATTGA AA 52

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SEQ ID NO:7709

SEQUENCE LENGTH:318

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08959

SEQUENCE DESCRIPTION:

GATCATACTN GTGATTATAA AAGNTCCTAG GAGGCTAGAA GAGCCAACCA ACAGAGAAGG 60
 GAAAGCAGTC TGTCTGAAC ATAGGGACAT AAGTTCATTG ATGCCAAGTA TCTTCCAGC 120
 ATGTTTCTCC CATTTAGAAT ATCTAGCATN TAAGGCCTTT CAATATTAAT ATAAGCCCAA 180
 TATCAGCTCT TTCTCTNTGT ATTTCATCTC TTTCTACTCT CCNATTTGTA TTTGGTGTTG 240
 CTNTTGAAAG TGTCGTATCT GGGAGATGAC CTGCCTTATC CTGTNCTATA ACAGTTTNTG 300
 NTTGCTGCTG TGTCTTTN 318

SEQ ID NO:7710

SEQUENCE LENGTH:243

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08960

SEQUENCE DESCRIPTION:

GATCCAAGAA ATTACTGAAA TGGACAAGAA ATGTCATTAG TTAAAAGCCA TGTTATCTTG 60
 AGCACACATA CAGTTAATTT GATAAAGAAG AGTATGCCTC TCAGCATTGT GAGATATATT 120
 TCAGACTAAT TTGATGTTCT TTAATGTTCC CTTATTTTGG AAACACGTTT TAAATGTGAA 180
 ATTAATGTCC TTCACAAAAT GTCTTTTGAA AGCACTGTGA CAGTTGTATA AAGATGATGT 240
 AAA 243

SEQ ID NO:7711

SEQUENCE LENGTH:122

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08961

SEQUENCE DESCRIPTION:

GATCTTGGCT TCTGAGGGCT CATTTGTAGA GTGGTGAGAA TAATATCTAA CTGCTAATC 60
 TTTCAGAGAC ATTCTGAGAG ACATATGCAA AAGAACTTG TAAATAAAAA GCACTGTACA 120
 AA 122

SEQ ID NO:7712

SEQUENCE LENGTH:240

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08962

SEQUENCE DESCRIPTION:

GATCCTTTAT TAAGCTTCTG ATTCATAAAA AAACCTATAA GAGACATTTG GGGGGATATT 60
 TGGGGAAATA GGAAGATGGA ATGGGTAGTA GATAATGTGG TATATTTNAT TTNCTAGAA 120
 GTGATAATTG AGTTACGTAG GATAATGTTT TAATTCCTAG GAGATGCATG CAGAAGTTTT 180
 TTTTTTAAAGT TTGAAAAGN CATGATGTTG NCAGCATATT TNCAAATAAN TATGTCCAAA 240

SEQ ID NO:7713
 SEQUENCE LENGTH:110
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08963
 SEQUENCE DESCRIPTION:
 GATCTGTGTC CAAAAGTGAA CTTNAGTCAG GAATGAATCA ATTCAGCAT AAACAAGCAC 60
 AAAANTTTAG TCTGCTGGCT GACTGGAAGC AAAAAAGTCA AGATGGNNNN 110

SEQ ID NO:7714
 SEQUENCE LENGTH:111
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08964
 SEQUENCE DESCRIPTION:
 GATCTCAAAA CAATTGTTGC AGCAGGCTCC TGGCAGTCTC AAGCAGTTCA TCTTCTTGGT 60
 GTACTGGTTT CCTATTGTGA TTTTATCATG GAAAATCAAT TGGCTNNNNN N 111

SEQ ID NO:7715
 SEQUENCE LENGTH:309
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08965
 SEQUENCE DESCRIPTION:
 GATCCTATAG GCAGAGAGTT TTCCTTCTG ACTTTTCCC TTTGCTTTCG TGTGACCACA 60
 TGTTTTCTGT ACCAGTCACT GGGGAAAGAA GTGAGTTTAT CTCGTTTGTT TAAAAAGTTT 120
 TGCTTGCTA TTTAGCATT CTTTTGGGT CTCAAGATT ATGGAACAAT AAATGTCATT 180
 TAATGCTGTG TGCTTATTTT GAATTCCTCA TCAGGTTTGA GAAGCGGGGT AAAAATACTT 240
 AGATGCTTAT CAGACTTGAA ATTATACTGA GTGGCATTGA ACGTGAGTTT GTCCCAGTGA 300
 ACCAGGCTN 309

SEQ ID NO:7716
 SEQUENCE LENGTH:204
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08966
 SEQUENCE DESCRIPTION:
 GATCCATGAA GCCAGAGCCT GGCTCTGGGT GACAGCAGGA AACGGCTGGG GAAAAGNACA 60
 GCAGGGCGTC CAGCTGAGCC TGCCATTCCA GCCTGTTTCGG GGGAAACAGC TCCCAATCCT 120
 AGACTGTGCT GCTCCGGGAG GGCTGCTGGT GGATATTGG GCTGTTTCA CCATATAGCT 180
 GTCATGAAAA ATGCTTCCGT GAAA 204

SEQ ID NO:7717
 SEQUENCE LENGTH:121

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08967

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SEQUENCE DESCRIPTION:

GATCTCAGGA ATACAGTCCC ATGCAAAGAT TCTCTGGTTT TATGGCTTTT TTCCCTTTCT 60
TTACACCATC CTCTCCCAATA AGCACCCATG TTTTTGAATA TGAATGTATT TGTAATAATA 120
A 121

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SEQ ID NO:7718

SEQUENCE LENGTH:186

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

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CLONE:HUMGS08968

SEQUENCE DESCRIPTION:

GATCACCTGA GGTCTGGGACT AAAAATACAA AAATNAGCTG GGCATGGTAG CCGGTNCCTG 60
TAATCCCAGC TATTTNGGAA ACTNAGAACA AGAGACTTGC TTGACCCTGG GAGGTGAAGG 120
TTGCAGAGAG CCGAGGTAGT GCCACTCCAC TCCAGCCTGG NCGACAGAGT GAAACTCCGT 180
CTCAAA 186

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SEQ ID NO:7719

SEQUENCE LENGTH:351

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08969

SEQUENCE DESCRIPTION:

GATCAGCTCT CTGTTTGGGA AAATGGAGTA GATATAGTTC TAGGTACTCC GGAAGACTA 60
GATGACTTGG TGTCAACTGG AAAGCTGAAC TTATCTCAAG TTAGATTCTT GGTCTGGAT 120
GAAGCTGATG GGCTTCTTTC TCAAGGTTAT TCTGATTTTA TAAATAGGAT GCACAATCAG 180
ATTCTCAGG TTACCTCTGA TGGAAAAAGA CTCAGGTGA TTGTTTGCTC TGCCACTTTG 240
CATTCTNTCG ATGTAAAGAA ACTNTCCGAG AAGATAATGC ATTTCCCTAC ATGGGTGAC 300
TTAAAAGGAG AAGGCTCTGT TTCCAGATTN CTGTACACCA TGTGTNAA N 351

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SEQ ID NO:7720

SEQUENCE LENGTH:142

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SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08970

SEQUENCE DESCRIPTION:

GATCCCAAAG AAGTAACGGG AAAAAGGAAT GTAAGGCATC TCGATAATNT TTTCATATTG 60
ATTACATGTT GAAATANTAT TTTGTATNTA TTGAGTTAAG AAAAATATGT CATTAAAAAT 120
AATNTCACCT GTTCTTTTA AA 142

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SEQ ID NO:7721

SEQUENCE LENGTH:130

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SEQUENCE TYPE:nucleic acid

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EP 0 679 716 A1

TOPOLOGY:linear
 CLONE:HUMGS08971
 SEQUENCE DESCRIPTION:
 5 GATCTTGTGA ATCAAATTCA CAAACATCAG CTTGTTACAA ATCATTNTAT TTATACCCCTT 60
 TAAAGAGTGT GGATTTAAGA TTCTTGAAAG GATGTAGNCT ATATANTTTC TCATTCCNTA 120
 TGGTGTAACN 130

10 SEQ ID NO:7722
 SEQUENCE LENGTH:180
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08972
 15 SEQUENCE DESCRIPTION:
 GATCGTTAAT TAATATCTTT CCATNCGAAG AGNAAGCAGA TTTCTTCTGA GATTTCTGAA 60
 CAGTTCCTGG GTGTTCCGG TTCTGCTTCA GTGTACTATC ACTTTGCCTT GCCTCAGTCT 120
 CTCCTTCCTC CGGGGCCACA GAAGGAGAAA GCCCAAAGCA TACCCATCCT GCCCTTTAAA 180

20 SEQ ID NO:7723
 SEQUENCE LENGTH:266
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 25 CLONE:HUMGS08973
 SEQUENCE DESCRIPTION:
 GATCACTGTC CCGNTTCAAA TTATCTTCA GTCCATTTC CCGGCCTATT TCAGCTGTTT 60
 CTTTTCACCT AACTGTTNAG TCATTNCGT TTTCAAGCAG TGCTTNATCT CATGTCCTTG 120
 AATATAGTTG TGTACTTTAT TTTNAGGTA ATAATTAGAA CAGTTCCTT CAGAGGCTGC 180
 30 ATTTGCCTTC TTCTGCCACC TAAATATTAC TTCCCTTCAA ATCNGCCTTT GAATCATCNN 240
 NTTTAAAAAA AANTTANCAT GNTTTN 266

35 SEQ ID NO:7724
 SEQUENCE LENGTH:33
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08974
 SEQUENCE DESCRIPTION:
 40 GATCAAAGGT ACCATGACCA AATAAGAAAC AAA 33

45 SEQ ID NO:7725
 SEQUENCE LENGTH:219
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS08975
 SEQUENCE DESCRIPTION:
 50 GATCTTCTTT TTTNTTCTGG AGCCAGACTT NCTGGGCTTN NATTGCTGCT ATGCTACTTA 60
 CTTGGAAATC TTGGACAAGT TACCTAATTT CTTGTTCTT AAGTTACTTC TGTTGTAATA 120

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TGATGTGGCC AATAATAGTG CCTATCTCAT AACATGGTTA TTAGAATTAA ACAAGTTAAT 180
ACTTTTAAAA TGCTTAGAAC AATTGTGGGT ACATACTAN 219

5 SEQ ID NO:7726
SEQUENCE LENGTH:275
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS08976
SEQUENCE DESCRIPTION:
GATCTGTCTG GNAACAGTTT ATGGAAATAT AGATATTCAT GCATCANATA AAAGTGCTGT 60
GACCATAGAT AAAGTGCAGG GAAGTTCTGT NACTGTATCT ACCGAAGATG GTTTGCTGAA 120
AGCCAAGTAT CTTTATACAG AATCATCATT TCTGTCTTCT GCTGCTGGGG ATATTACATT 180
15 AGGAAAGTGT CATGGNAATA TAACATTACA AAGCAAGANG GGTAGCATCA CAGTNGATTG 240
GTCTTCTGGA TGTCTAAAAG CCTCAACTAA TCAGN 275

20 SEQ ID NO:7727
SEQUENCE LENGTH:37
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08977
SEQUENCE DESCRIPTION:
25 GATCCCAGGT TCTCATGAAC CAAATAAGTA ACTTAAA 37

30 SEQ ID NO:7728
SEQUENCE LENGTH:97
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08978
SEQUENCE DESCRIPTION:
GATCTGCCCC CCTNGGCCTC CCAGAGTGCT GGGATTACAG GCATGAGCCA CTGCGCCCGG 60
35 NCTGTACTAA GTCITTTTTT TTTAAATTC CTCNAAA 97

40 SEQ ID NO:7729
SEQUENCE LENGTH:284
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08979
SEQUENCE DESCRIPTION:
GATCTATGAA CTTCTGCCCC ACATAACTTT TAAAGCAGTA TTTTATAAAC CCCATGAAGA 60
45 GCTTGCTTTT TTGTCTTTGT AATTCCTTAG GGTCTGTGAT ATATGCCAGG AAATCTGATT 120
GAAACTAATA TGCTTTTTAT TCTCCCATTA TTCCCTAAAT NGTTATATCA CAGGCACTTG 180
CCTACATTGG GAAAGTAAGG ACAAATAATA CCCATTTAAA GTAAGTNGCC TATTATTNNG 240
ACTGTCTTTC GTAAGCACTT GCTGAACNGC TGACTTAACT CAGN 284

50 SEQ ID NO:7730

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EP 0 679 716 A1

SEQUENCE LENGTH:374

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08980

SEQUENCE DESCRIPTION:

GATCTCGCAA TGGGGAAGGC CCTGGTCGAC CAGAGGGAGC TGTACCTGGG CCTGCTCTAC 60
 CCCACGGAGG ACTACAAGGT ATACGGGTAC GTCACCAACT CCAAGGTGAA GTTTGTCATG 120
 GTGGTAGATT CCTCCAACAC AGCCCTTCGA GACAACGAAA TTCGCAGCAT NTTCCGGAAG 180
 CTACACAACT CCTACACAGA CGTGATGTGC AACCCCTTCT ACAACCCGGG GGACCGTATC 240
 CAGTCCAGGG CCTTTGATAA CATGGTGACG TCGATGATTG ATACAGGTTG TGCTGAGTGA 300
 GCTGTGCTGC CAGCCATCGN AGAGGAGCCC NTNGCACGAC TTGTGGTGGG GTCCGTCGGT 360
 CTTGTTTCTN TTTN 374

SEQ ID NO:7731

SEQUENCE LENGTH:164

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08981

SEQUENCE DESCRIPTION:

GATCCGGTTG GACTCTGACA TCGGATGCCC TCAAACATAC AGAACTTCCA AACTCAAGTC 60
 CAGCCATAAG CTATTTTGCC AACATGTCAG AGTAATCTGT ATTTTGTAT GTGATTCTA 120
 CTTTATAGA CTTGTTTAA AACATAAAA CACATTTTAA TAAA 164

SEQ ID NO:7732

SEQUENCE LENGTH:287

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08982

SEQUENCE DESCRIPTION:

GATCTTTCAT GAAAATTAAT TGTGCCCATG TCCAAGTTTG AATTAGAGAT ACACAGCACA 60
 CAATCATTTT TGTTACCACT TTTGGAATAT CTAGCATTAG CCTTGATAGT TTTGTGTGGT 120
 GTGTTTTGAG TATATCTGAN CTGTTAGTTA TATTTGGTTA ATTTATTTAA NGATGTGTGT 180
 TAANCCTTAA TATTTATGCA GTGTTTAAGT ATTTGGAATA TATTNGAAAT AAATTATCCA 240
 GTGTTTTAGA TAAATGGTTA TTAAATTGCN NAANGTGATT CTCTAAA 287

SEQ ID NO:7733

SEQUENCE LENGTH:173

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS08983

SEQUENCE DESCRIPTION:

GATCTGTGCC GTTCAATAGG CATCGTCTCT CAGCCTGAGG GAGGCTGGAT TCTGGGTTCC 60
 TGTAATCACA GGGAGGAAAA GCTTTCTTAA AAATGGACAT GTATGTGCGT GTNAGTGTGT 120
 GTGTAGATTT ATAGTTTTTG GTAGTGGCAG GAATAAAAAA AATCCATCCT AAA 173

EP 0 679 716 A1

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SEQ ID NO:7734
SEQUENCE LENGTH:198
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08984
SEQUENCE DESCRIPTION:
GATCTCTATT TTGTGCCAG ACTCTGAGCT GCTTTATGAA CACCTCAATT AAATNACTTC 60
ATTGGAGTGA TTTGAGTAAT CTATCTATAG TGTAATATTT TCTATGAATA ATTTTCTGTA 120
ACCTTATTTT ATAAGTTAAT ATCTAAAAAG TGTTTATTTT CAAATTGCCA AATAAAAGGT 180
TTTATATTAG TCATTAAA 198

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SEQ ID NO:7735
SEQUENCE LENGTH:94
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08985
SEQUENCE DESCRIPTION:
GATCTAGAGA CTCATTCAAT AGCAATGTGA CCTTTTAAAT ACTTACATTA AGTAAACTG 60
CCAGTAGATT AAATCATATA TATATATATA TATN 94

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SEQ ID NO:7736
SEQUENCE LENGTH:363
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08986
SEQUENCE DESCRIPTION:
GATCACCTGN ACTCCAGCTC AGATTGCCCTC TCCTGGACAT GGCAATGAAT GAGTTTTTAA 60
AAAACAGTGT GGATGATGAT ATGCTNTTNT GAGCAAGCAA AAGCAGAAAC GTGAAGCCGT 120
GATACAAATT GGTGAACAAA AAATGCCCAA GGCTTCTCAT GTCCTTATTC TGAAGAGCTT 180
TAATATATAC TCTATGTAGT TTNNTAAGCA CTGTACGTAG AAGGCCTTAG GTGTTGCATG 240
TCTATGCTTG TGGAACTTTT CCAAAATGTGT GTGTCTGCAT GTNTGTTTGT ACATAGAAGT 300
CATAGATGCA GANGTGGTTC TNCTGGTACG ATTTGATTCC TGTTGGGATG TTTTAACTAA 360
NAN 363

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SEQ ID NO:7737
SEQUENCE LENGTH:262
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08987
SEQUENCE DESCRIPTION:
GATCTTCTTT GATGATGAGA GGCAGGAATAT TGTAAGACGTC AGCAAACTGG GTGTTACCTG 60
CATTCACATC CAGAATGGAA TGAATCTTCA AACTCTAAGT CAAGGGTTAG AGACATTTGC 120
AAAGGCCCAA ACTGGGCCTT TAAGGTCCAG CCTTGAGGAG AGCCCATTTN AGGCCTAAAC 180
TGAAAGGAAA TCAAGAAGGC ATTTTCAGGT GCATTTGTAA TTTATTAAAG TTCATCTGTG 240
TGTGACAGAA AAAAAAAGTA AA 262

EP 0 679 716 A1

SEQ ID NO:7738
SEQUENCE LENGTH:361
SEQUENCE TYPE:nucleic acid
5 TOPOLOGY:linear
CLONE:HUMGS08988
SEQUENCE DESCRIPTION:
GATCTGTGCC ACATGGAGGT TTTGGGATGG GATTNAACG CTACCTGCAG TGCATCTTGG 60
GTGTTGACAA TATCANAGAT GTTATCCCTT TCCCAAGGTT TCCTCATTCA TGCCTTTNAT 120
10 AGCTGGAAGA TTGGTTAAGG CAAAGCACCC CCCATGGCAG AGACACTGCA CATGATTGTN 180
CATACAGCAG AATGCATGTT TGGATTNTAG AAATGCAGAT TTCAATATGT AATTGTTGTG 240
CCATAAGATA TCATAGNAAA NNTATAAGTG GTTGTNCTTT TCTTAGAAA TTAGAGGGTAT 300
TTCACGTAAG GATGAGCTCC CGCAAGNNGA GGTACTTNTT AGCANGGGGA CTCTCAATCG 360
15 N 361

SEQ ID NO:7739
SEQUENCE LENGTH:143
SEQUENCE TYPE:nucleic acid
20 TOPOLOGY:linear
CLONE:HUMGS08989
SEQUENCE DESCRIPTION:
GATCTTGATG TGGGTGATAG TTTTACTTAT GTACATCGAT ATACATCTAT TTAAAAACAT 60
25 GTACATTAA ATTGTTGCAT TTTATGTATT TAAAATTAT TGAGTANTCC TTCTGTATGT 120
ATTATTAAAG AGAATATAGC AAA 143

SEQ ID NO:7740
SEQUENCE LENGTH:333
30 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08990
SEQUENCE DESCRIPTION:
GATCATAGAG TTTGGCTGGG GAGGGGGGCA GTTTTAGAGG CTTCCACTTG GTGTTCTCA 60
GAATGATATC TCTTACTCCG GGGGCCAAGG TAGGGGTTAG CTTTGTCTCT CTTTGTAGTT 120
TAGATTGTAT CTCTGCCTT GTTCAAGTTC ACAAATCTTT TTGTGTATAC ACATATGTAC 180
ATGAAAAATNA TGTTTCATGCT TTTAATTATT TTACCCCTCA TTATTTTATT TTTNATAGTT 240
CTCATAGCTA TGTCTTTCAG TTCACTAATC TTTTTCAC AGTGTTTAA CTGTCATTAA 300
40 ATCCCATCCA AAGTATGTTT TCTCTCAGAA GAN 333

SEQ ID NO:7741
SEQUENCE LENGTH:135
45 SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08991
SEQUENCE DESCRIPTION:
GATCCTTGTA CACTGTTTAT ATGTGCAATA AAATGCATGG CCAGCTGACA GTATCGTCAG 60
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TGGACAGAAT GTATATAGTG CAAATATCCT TCCTTTTATT ATTTAACAGC CATTAAAATC 120
TTGGATTTGT ACAA 135

5 SEQ ID NO:7742
SEQUENCE LENGTH:333
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS08992
SEQUENCE DESCRIPTION:
GATCCCAGTT CACAATTATT CACCGGGTAT TTAGTTATTC ANAGGCTGCT CTGCTGAGAA 60
GNTGAACAAA TTTCTTGTC AAAACAATGT ATTTCAAACG TGCCGCTCGG GCCTTTCCCG 120
TATTGCTCAC TGGTGCTGGG AAGACAACAC TTCTGAACTA TATTTTGACA GAGCAACATA 180
15 GTAAAGAGT AGCGNNNNCT TTAAATGAAT TTGGGAAGG AAGTGCGCTG GAGAAATCCT 240
TAGCTGTCAG CCAAGGTGGA GAGCTCTATG AAGAGTGGCT GGAACCTAGA AACGGTTGCC 300
TCTGCTGTTT AGTGAAGTGA GGAATGTGTT TTN 333

20 SEQ ID NO:7743
SEQUENCE LENGTH:219
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08993
25 SEQUENCE DESCRIPTION:
GATCCCACT GTATAACATT CTGAAAAAGG CACAACTATG GAGACAATAA AAAGAACAGA 60
GTTTTGGATG AAGGGAGGGA TGAACAGGCA GAATACAAAC AAATTTTAGG GCAGTGAAAC 120
TATTATTATG ATACTTTATG GATATAGAAT ATTATGCATT TTTCAAAACA CATAGAATAT 180
30 ACAACATGAA GAATGAATGC TAATGTAAAC TGGATTTTN 219

SEQ ID NO:7744
SEQUENCE LENGTH:59
SEQUENCE TYPE:nucleic acid
35 TOPOLOGY:linear
CLONE:HUMGS08994
SEQUENCE DESCRIPTION:
GATCTTGGGC CACACATGAA AATACTAA CACTAATGAC AGCTGATGAG ATGCTTAA 59

40 SEQ ID NO:7745
SEQUENCE LENGTH:353
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
45 CLONE:HUMGS08995
SEQUENCE DESCRIPTION:
GATCAGAGTG ATGCTGGAAC TCAGTGTGCA TAAAATTTCA GTCAGTGAAT ATCACTGAAC 60
GTCATATACT ACTTGGTATG TGACTTTGGT TTGTGTTAAG AAAGCTTGTA TATAATATT 120
50 TTNGCCATAG TAAGTGAGAA ATTGCTTA ATCATGCCTG TTTGATGGTA CTAGGAAAGA 180
AAGGGGTAGA GATTAATTCT TGCACAGTAT AAGCAACAGT GCAACAACT ATGCCATTTN 240

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EP 0 679 716 A1

CCTTTTNCCT CTTACTTGAA GGCAGAATCG CAAAACGTTT GANATGGCTT TCCTAAACTA 300
CTCTACTCTG GTGNGAGCTC ATTTACCACA NGAAGCCTTA TAACACCGTT TNN 353

5 SEQ ID NO:7746
SEQUENCE LENGTH:327
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS08996
SEQUENCE DESCRIPTION:
GATCAGGTGA TTGCCTTTCT CAGCTGTCAG TTCTCTAATT TCAGGCTTGG TAGCTTGTAG 60
GAACTGAAAT TGCAATTAAA ACCTTTATAA ACTCAAATA AATCATGAAT TACAGAAAAA 120
GTCCATTCTT CCAAACTTG ATGTTACCAC ACTTACAAGT TTAATAATG AAGTCGACTG 180
15 TTTAAAGGAT TCTGCATATA TTCTAGTGTG CACATTCAGA AACATTTTTC TTGGAAAAAG 240
TACCCAACAT TTTTATAAC TGCACATATT AATTTATTGC CAGAATAAAT TGCATTGCAT 300
GCTAAATTAA NGTCAGATAN TTCNAAA 327

20 SEQ ID NO:7747
SEQUENCE LENGTH:153
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08997
25 SEQUENCE DESCRIPTION:
GATCATGTAA ACTTGCTGTT TTNGTTTTN CCTGCCGGGT GTTGATGTG TGGTGACTTG 60
CGGATTTATG TTTCAGTGTG CTGGAACTT TCCATTTTAT TCAAGAAATC TGTTCATGTT 120
AAAAGCCTTG ATTAAGAGG AAGTTTTTAT AAA 153

30 SEQ ID NO:7748
SEQUENCE LENGTH:41
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
35 CLONE:HUMGS08998
SEQUENCE DESCRIPTION:
GATCTTAATT TTGTTTTTGG TTAAATAG TGTTTCCTTT N 41

40 SEQ ID NO:7749
SEQUENCE LENGTH:299
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS08999
45 SEQUENCE DESCRIPTION:
GATCTATCTA AATGCCGATT TGAGTTCGCG AACTATGTA CTGCGTTTTT AATTCTTGTA 60
TTNANACTAT TTAATCCTTT CTACTTGTCG CTAAATATAA TTGTTTTAGT CTTATGGCAT 120
GATGATAGCA TATGTGTTCA GGTATATAGC TGTTGTGTTT AAAANTTGAA AAAAGTGGAA 180
50 ANCATCTTTG TNCATTTAAG TCTGTATTAT AATAAGCAAA ANGNTTGTNT GTATGTNTGT 240
TTNATATAAC ATGACAGGCA CTAGGACGTC TGCCTTTTTN NGGCAGTNCC GTTAAGGGN 299

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EP 0 679 716 A1

SEQ ID NO:7750

SEQUENCE LENGTH:169

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09000

SEQUENCE DESCRIPTION:

GATCAATTAG ACATTTTGAA AATAATTAA AGTGTTTCC TTAATGTTCT CTGAAAACAA 60
GTTTCTTTG TAGTTTAAAC CAAAAAGTG CCCTTTTGT CACTGGATTG TCCTAGCATT 120
CATGATTTT TTTTCATACA ATGAATTAAA ATTGCTAAAA TCATGGAAA 169

SEQ ID NO:7751

SEQUENCE LENGTH:53

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09001

SEQUENCE DESCRIPTION:

GATCATTTAA TGAATCCTCA AGGACTAATG AAATAAATGC TAGACTGCTG AAA 53

SEQ ID NO:7752

SEQUENCE LENGTH:252

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09002

SEQUENCE DESCRIPTION:

GATCCTGAAA ATAAAACTAA CACTCCAGTA TTTTGTCTT GTTTTTCGCA ATTGAGCTAT 60
CTGAAAAGTG TTATTCCTAA GTAATGTTCA AAAATGATAA GTAATCTGGA TACCTTTTTC 120
TTATACTTTC TCCTAGGAAA ACTTTAAAC TTTAAAAAGG CAAACCTACC AATAGGAATA 180
ACANATTAAA TGCAAGAGA GTATATCCAA TATTAGGATA TAANTGTATG TGTCTCAAGT 240
TTAACTCTAC AN 252

SEQ ID NO:7753

SEQUENCE LENGTH:237

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09003

SEQUENCE DESCRIPTION:

GATCTACAAA AAATTCCTGG AGCCATATAT ATACCCTCTG GTNTCCCAT TCGCTAGTCG 60
TATATNGCCT NAGAAAGCAA TACAAGAATC CAATAATACA AACAAAGGCA NAGTAACTT 120
TAAGGGNGCA GACATGAATG GCTTACCAAC AAAAGGACCA NCAGAAATCT GTGATAAAAA 180
GAAAGACTAA AGAAATTTTC CTAAAGGACC CCATCATTAA AAAANTGGNC CNGGNNN 237

SEQ ID NO:7754

SEQUENCE LENGTH:264

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09004

SEQUENCE DESCRIPTION:

5 GATCTTCCCA AACCCAGAGA AAGACATCAA TTTCCAAGAA CAAGAAGGTT ATCAAACACC 60
 AGGCAGATTT AACCCAGAGG ATACTACCTC AAGGCATTTA ATAATCAAAC NNCCAAAAGT 120
 CAAGAATAAG GAAATAATCC TAAAAGCAGC AAGAGAAAAG AAACAANTAA TATACANTGG 180
 AGTTCCACTA TATCTGGCAG CAGACTCTTC ATTGGAAACC TTATAGGCCA GGNGNGAGTG 240
 10 GCATGACACA TTAAANGTAC TGGN 264

SEQ ID NO:7755

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

15 TOPOLOGY:linear

CLONE:HUMGS09005

SEQUENCE DESCRIPTION:

20 GATCCAGAGC TGTTGCCTGT GACAGCGGTT TCTNTGGATG TCAAAGGCAG CTGCCTGGTT 60
 GCCCAGCTTG CTTCTCGACT GGTGGCCCTT ATGGGTGGGT GTGCGATGGA AATNTGTTCC 120
 TGCCGGAGTC TGAGGCACCA GGGTGTGCTC AAAGGCTGGC CCTGGTGGTG GACTGGCACC 180
 TGTGCAGAGT GCCGTGTGCT TGTGGTGC GC CATCTGAAGC AAGAGTCCAG CGTTCTGCCG 240
 TGCTGTCCC CCACCATGCC CCCTACAGGC GGTACTGATG GCGCTN 286

25 SEQ ID NO:7756

SEQUENCE LENGTH:382

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09006

30 SEQUENCE DESCRIPTION:

35 GATCCTTTTT CCTTCAGCAG CATTTCTTAC TGGCTGTGGC TGGAATCTGC CTTTAATCAC 60
 AGCTGTCAAC ATTCTCAGT GATTCTTGTG AGACTCTTTT TGGTTATAAT TACTATTTAA 120
 TATTTAGACT ATTTTACTGA GCAGACTTTA TAAATGAGAT ATCTACAAGG CACTTAAAGT 180
 GTTACAGATG TTTTACCTTA AGANTTATTT AAGTNGTGTT GGGTTAAGAC AGTTTTCAGT 240
 GTACCGTAAA TGTGTGTTT TCAGAAAAAG ACAAACGAT GGTGCTGACT GGTTTTCTGT 300
 ATATNGCACA ACAGTCTCA AATACACTGG ATGTATGGAA CCTNTTCCTT NCATCCAGGC 360
 AGCATTTTTN TTNCNCTCTN CN 382

40 SEQ ID NO:7757

SEQUENCE LENGTH:298

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

45 CLONE:HUMGS09007

SEQUENCE DESCRIPTION:

50 GATCATCTTT GAGTAGCACT GTTTGGGGC CCTCGGTCTC TCTGAAGACC CTAGCAGAAC 60
 TGATACCTAC CTGTATCTCT TGTCTCTCC TATTTGAGTT TCACTTCCAG AGAACTTGTT 120
 CTTCAGCAAG AATGTGTCAC TAGTAAGGAC ATCTCTAGCA TTTCTTAGC CTTCTTTTC 180
 TGCTGCTCAA AAATAATCGT TACAAAGCTT AGGTTTAAGC TGTATATGAA ATATTTATGC 240

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GACTCTCAAA CTTTAAAGGA GTTGCTCCTT TGTTCACAAA TTAAATGTGT TAGATAAA 298

SEQ ID NO:7758

SEQUENCE LENGTH:292

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09008

SEQUENCE DESCRIPTION:

GATCATGGTC TTCAAAGATG TAAATACAAG AGAACCAAAG TTATTCCGGG CAGTTCCAGA 60
 TTCAGAACGT GTTATTCCTC CCAGAACGAC AACGCCATAT TTCCTTTGAG GCCTTCGCCC 120
 ATCCTGCTGA CCCATTTTNN TGCCCTCTTC TTACCCCAAT TTTNTTGTAT TACCCTCTAC 180
 AATATACTTT TNATTGAGCA CTTTGCTGCT GAAATGCTGC CTCTTGCCTT TTTTTTTTNA 240
 AATTTTAAAT TATCTAAATT TATGGTTTGT GGTGGTGTCT ATNGCAAAGT TN 292

SEQ ID NO:7759

SEQUENCE LENGTH:307

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09009

SEQUENCE DESCRIPTION:

GATCTGTAA TGAAAAGTGC TGGAATACCT CAACCAAGAA AAGGCAGAAA AGGGAAGGGA 60
 GCATATTGCT AAGAGAGCAA TGGAAAATAT GTAAGCTGCT TTCATTAATT ACCCTACTTT 120
 CATTCTCTCC ACCCAAGCA AATCCCAACA TTTCTCTTCA GTGTGTTGAC TTCTATCCTG 180
 TTAACACTGT AATATCTTTA AATGATGTAC AGGCAGATGA AACCAGGTCA CTGGGGAGTC 240
 TGCTTCATTT CCTCTGAGCT GTTATCTTGT GTATGGATAT GTGTAAATGT NGGTGACTCC 300
 TTGATAN 307

SEQ ID NO:7760

SEQUENCE LENGTH:165

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09010

SEQUENCE DESCRIPTION:

GATCTCTGAA CATGAAAACC AAGGTGGCTA TTTTCAGGTT GCTTTCAGCT CCAAGTAGAA 60
 ATAACCAGAA TTGGCTTACA TTAAAGAAAC TGCATCTAGA AATAAGTCCT AAGATACTAT 120
 TTCTATGGCT CAAAAATAAA AGGAACCCAG ATTTCTTTCC CTAAG 165

SEQ ID NO:7761

SEQUENCE LENGTH:334

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09011

SEQUENCE DESCRIPTION:

GATCGATGAC GCAGGTCCTT CCAGTGAAAT GAATGGTCTC AAAAAATGGA GAGGATGTGC 60
 TGGGGATTGA GAAGGCAGGG TAGCCAGAAA GAAGCACAAA TTACCCCCAC CATAAGCTTC 120

EP 0 679 716 A1

ACAGAATTGC TGACGCTAAC GGCCATGGGT GAGATTTTGT CAATAAGTGT CAGCCTGATG 180
ACAGATACCA CAGACAGTTG CAAAAATGTC CAAGTTTCTC CTGAAGTTT CTGGCAGGGC 240
AGTGGGCTCC ATCCTTAGGA GTGAATCAGC TTTGGTGAGT ATCGTGATTG AACAGAGCCT 300
5 TAACACCAGG GCCCCCTTAA CTCCGCACTC CTTN 334

SEQ ID NO:7762
SEQUENCE LENGTH:247
SEQUENCE TYPE:nucleic acid
10 TOPOLOGY:linear
CLONE:HUMGS09012
SEQUENCE DESCRIPTION:
GATCGCTTTG ATTTTAAAGT TCATTGGAAC TACCAACTTG TTTCTAAAGA GCTATCTTAA 60
15 GACCAATATC TCTTTGTTTT TAAACAAAAG ATATTATTN GTGTATGAAT CTAAATCAAG 120
CCCATCTGTC ATTATGTTAC TGTCTTTTTT AATCATGTGG TTTGTATAT TAATAATTGT 180
NGACTTTCCT AGATTCACCT CCATATGTGA ATGTAAGCNC TTAACATATGT CTCTTTGTAA 240
TGTGTAN 247

20 SEQ ID NO:7763
SEQUENCE LENGTH:312
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
25 CLONE:HUMGS09013
SEQUENCE DESCRIPTION:
GATCCTTCCT GAGCGTTTCC TGGCAGTTTT AAAACGAAAA ATCAGTGTTA AGTTTGATGC 60
AGTTATTGGA TATAAATGA AAGCGCAATA AGCACCTAGT TTTCTGAAAA CTGATTACC 120
30 AGGTTTAGGT TGATGTCATC TAATAGTGCC AGAATTTTAA TGTGTGANT TCTGTTTTTT 180
CTAATTATCC CCATTCTTC AATATCATTT TTGAGGCTTT GGCAGTCTTC ATTTACTACC 240
ACTTGTCTT TAGCCAAAAG CTGATTACAT ATGATATAAA CAGAGAAAATA CCTTTAGAGG 300
TGACTTTAAG GN 312

35 SEQ ID NO:7764
SEQUENCE LENGTH:93
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
40 CLONE:HUMGS09014
SEQUENCE DESCRIPTION:
GATCTGCCTG TCCGGGCTGG GACAGAGACT CCCCAAGNAC CCCATTCTGC CTCCTTCTGG 60
GGAAATAAAT AAGTGTCTGT TTCAGCAGCT AAA 93

45 SEQ ID NO:7765
SEQUENCE LENGTH:133
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
50 CLONE:HUMGS09015
SEQUENCE DESCRIPTION:

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GATCTGGCTA AGCATCTTCC TCATGCTTTC AGTAAATTGG AAAACCCACG GAGGTAAAT 60
ATTACCTTTT TTTTAACTAA CTAAANCA TNCTGTTTAT TTTTNATTA CATTNCCCA 120
TATGNGNGGA ATN 133

SEQ ID NO:7766
SEQUENCE LENGTH:300
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09016
SEQUENCE DESCRIPTION:

GATCGTNAAA ACCCAAATG GACTTATTTT AAGTCACTTG CCAAATGGCC CAACTGCTCA 60
TTTAAATN AGCAGNGTTC GTCTTCGTAA AGAAATTAAG AGAAGAGGCA AGGACCCAC 120
AGANCACATA CCTGAANTAA TTCTGAATAA TTTTACANCA CGGCTGGGTC ATTCANTTGG 180
NCGTATGTTT GCATCTCTCT TTCCTCATAA TCCTCAATTN NTCGGNAGGC AGGTTGCCAC 240
ATTCCACAAT CAACGGGATT ACATATTCTT CAGATTTCAC AGATACATAT TCAGGTGTGN 300

SEQ ID NO:7767
SEQUENCE LENGTH:105
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09017
SEQUENCE DESCRIPTION:

GATCAAGTA GCCACAACAT TCTCTTAAGC CAAAGNCCAA TCCAGAACAA NGCCCTAACT 60
CTCTTCAATT CTATGAAGAC TAAAAAAGTA AGGAAGCTNC AGANN 105

SEQ ID NO:7768
SEQUENCE LENGTH:216
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09018
SEQUENCE DESCRIPTION:

GATCGTCAAA CCATTCATCC TTTTAAAGGT TTATTTGAAG ATGCTGTTAA AGTACAGAAT 60
TTTGTGTACA GGTAGATTTT TCCGTCCCTC ATTAATAGTG CCTTCTTAAT TAATACAGNC 120
TGGTGTAGC TATAACAAGG CNCCAGTAAG GCCAAAGAAT CCCAAGTTCT TTGTGGAAAA 180
ANAAAAAANN NCTTTTGGGG NCAGNTTTC CCTTCN 216

SEQ ID NO:7769
SEQUENCE LENGTH:75
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09019
SEQUENCE DESCRIPTION:

GATCCTCCCG TGTGTGGTTG GAAAACTTTT GTTTTTTGGG GTTTTTTTTT TCTGAATAAA 60
AAAGATNCTA CTAAT 75

SEQ ID NO:7770

SEQUENCE LENGTH:341

SEQUENCE TYPE:nucleic acid

5 TOPOLOGY:linear

CLONE:HUMGS09020

SEQUENCE DESCRIPTION:

10 GATCTGTGTC TTAATTGTTT AGTTAGAGTG AGAAGTTGAC CTATGATTCT TTTTAAATT 60
 TTATATTGG AACAAAGCTG CAAGTTATGG TAANGTACTG TACTGNGNGA NGTATTATGA 120
 TATTTAATGC ATCTNTGGCT TAACACTTGT GAGAGTTACC AGCTTGAAAA TGATGGTGTT 180
 GACTACCTCT TGANTCACAT CTATCAACCA CTGGCACCTA CCACCAAGCT GGCTTCAATT 240
 AGTATGTGTT GCTTTTGGT ATTAACANCT AACCGTACTA GAGACCAAAG TGAACCCTGA 300
 TTTTAAATAT GTNTTAAATA ATGGTGGTTT TATCTAGTGG N 341

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SEQ ID NO:7771

SEQUENCE LENGTH:238

SEQUENCE TYPE:nucleic acid

20 TOPOLOGY:linear

CLONE:HUMGS09021

SEQUENCE DESCRIPTION:

25 GATCCAAGCC TGGAGTTTGC AGAAGATACT GTCCTAATAA GCAGGCATT CTAAACCAAG 60
 TATCTAAGCC TAAGCACAGC TTGTCCTGGG TGAAATGTCT GCCACAAAAG ATAGTTTCTC 120
 CTAGCTCAGA CTTAACCATT TATAAAGGTT GGTAAAATAC TGGCAGTGAC AACAAATTGA 180
 CTTTTAATT TTTTATTTG CATTATTCCA ATAAATGAAA ATCTGTCAGA GTTCTAAA 238

30 SEQ ID NO:7772

SEQUENCE LENGTH:123

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09022

SEQUENCE DESCRIPTION:

35 GATCTGCCCC CACTGTCCTG TGAGGACAGC TGAGGCCAAG NAGTGAAAA CCTATTACTA 60
 CTAAGAGAAG GGGTGCAGAG TGTTACCTG GTGCTCTCAA CAGGACTTAA CATCAACAGG 120
 AAA 123

40 SEQ ID NO:7773

SEQUENCE LENGTH:361

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09023

45 SEQUENCE DESCRIPTION:

50 GATCCAGGTA GAATTTTNA ACCAATCAAA TGAAAAAAC AAACAAACAA AAAAGGAAAT 60
 GTCATGTGAG GTTAAACCAG TTTCATTCC CCTAATGTGG AAAAAGTAAG AGGACTACTC 120
 AGCACTGTTT GANGATTGCN TCTTCTACAG CTTCTGAGAA TTGTNTTATT TCACTTGCCA 180
 AGTGAAGGAC CCCCTCCCCA ACATGCCCA GCCCACCCT AAGCATGGTC CCTTGTACC 240
 AGGCAACCAG GAACTGCTA CTTGTGNCC TCACCAGAGA CCAGGAGGGT TTGGTTAGCT 300

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EP 0 679 716 A1

CACAGGACTT CCCCCACCCC AGANGNTTAG CATCCCATAC TAGACTCNTA CTCAACTTNN 360
N 361

5 SEQ ID NO:7774
SEQUENCE LENGTH:54
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
10 CLONE:HUMGS09024
SEQUENCE DESCRIPTION:
GATCAAGCGA ACTAATTTTA GGTGAAATC CGAATAAAAG AACTTTACTG GAAA 54

15 SEQ ID NO:7775
SEQUENCE LENGTH:188
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09025
20 SEQUENCE DESCRIPTION:
GATCTGTATA ATGTACATTC AATATAGAAA GCTTTATATA TTAAAAAGNG TATAGAACAT 60
TTNACAATTA CACTCATCTT TAACATAACA TCTTGACATC CATTTTNAAG TTTTTTTGCA 120
CAAGCTCCTT TNCATTCAAT TTGGTAAAGC CAGTTATACA TACTANTGTG TACTGTGAGC 180
TTTCAGAN 188

25 SEQ ID NO:7776
SEQUENCE LENGTH:221
SEQUENCE TYPE:nucleic acid
30 TOPOLOGY:linear
CLONE:HUMGS09026
SEQUENCE DESCRIPTION:
GATCACCTGN AGGTCAGGAA GTTTGAAAAC CAGCCTGGCC AACATGGCAA AACCCTGTCT 60
CTACTAAAAC AAAAATNAGC CAGCATGATG GCAGGCGCCT GTAGTCCNNN CTATTCAGGA 120
35 GGCTGAGGCA GGAGAATCGC TTGAACTGGG AGAGGGGCAG AGGTTGCAGT GAGCTGAGAC 180
TGCATTCCAG CCTGGGCGAC ACAGCAAGAG TCTGTCTCAA A 221

40 SEQ ID NO:7777
SEQUENCE LENGTH:200
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09027
45 SEQUENCE DESCRIPTION:
GATCGTAGGA ATTGAAGGAG TGTCCCGCCT TGTGGCTGAN AACTGGACAG TGGCAGGGGC 60
TGGAGATGGG TGTNTGTGTG TGTGTGTGTG TGTGTGTGTG TGTGTGCGCG CGCGNCAGTA 120
CAAGACCGAG ATTGAGGNA A NCATGTCTT CTGGGTGTNA CCATGTTTCC TCTCAATAAA 180
GTNCCCCTGT GACACTCAAA 200

50 SEQ ID NO:7778

55

EP 0 679 716 A1

SEQUENCE LENGTH:286

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09028

SEQUENCE DESCRIPTION:

GATCCTGGTC CCCTCCCTGC TGGACCTGCC ACCCAGAGCT TCCTGNCTAG TCCCACTGGG 60
CTGGCCACC AGGCCTCTGA CCCAGGCTGC TCTNCGGCC CTTCCTCCTC CTNTTCCTGC 120
TCCAATTCT GTCCACCTGG GGACAGTCTG TGCTGTAGC CTNATGACCC CAACCCAGCC 180
CCAGGCATGG CTAACCCCTG ACTGNTTGCC TCATATTTAA GCTGCTGNTN TGGCCAAGTG 240
NCTAATTTTA ACCCAGACCT NAATAAAGAC ACNTNTTGT ACCAAA 286

SEQ ID NO:7779

SEQUENCE LENGTH:127

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09029

SEQUENCE DESCRIPTION:

GATCTGATTT TTCATAAAAA ACATTGTGA ACCTTCGGCA TAAATGGGT AAGGTGCCAT 60
CCCTGAAACT GCAATGCAGA TATGTTCAAG TAACCTTAT TTTTAATTA AAAATAAATC 120
TTTCAAA 127

SEQ ID NO:7780

SEQUENCE LENGTH:105

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09030

SEQUENCE DESCRIPTION:

GATCTGANTT TTCATAAAAA ACATAAANTG ACCTTCGGCA TAAATAGGT AAGAAGCCAT 60
CCCANAACT NCAATGCAGA TATGTTCAAG TAACCTTTAT TTTTN 105

SEQ ID NO:7781

SEQUENCE LENGTH:321

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09031

SEQUENCE DESCRIPTION:

GATCGGCGCC GTTACAGACG TGGCTACTAT GGAAGGCGCC GTGGCCCTCC CCGGAATNCT 60
GGTGAGATTG GAGAGATGAA GGATGGAGTC CCAGAGGGAG CACAANTNNN GGGACCGGTT 120
CATCGAAATC CAACTTACCG CCCAAGGTAC CGTAGCAGGG GACCTCCTCG CCCACGACCT 180
GCCCCAGCAG TTGGAGAGGC TGAAGATAAA GAAAATCAGC AAGCCACCAG TGGTCCAAAC 240
CAGCCGTCTG TTCGCCGTGG ATACGGCGT CCCTACAATT ACCGGCGTCG NCCGNTTCCT 300
CCTAACGGTN CTTACANGT N 321

SEQ ID NO:7782

SEQUENCE LENGTH:66

EP 0 679 716 A1

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09032

SEQUENCE DESCRIPTION:

GATCCACTGA GTTAGTTTCA TTGGGGCGGG GGAAAGAACT GGAATTAAAC TTGTTTAATC 60
CTTAAA 66

SEQ ID NO:7783

SEQUENCE LENGTH:131

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09033

SEQUENCE DESCRIPTION:

GATCCTCCCA GCTGTCGGCC GCGGACCCGG GCCGCGTGTG AGCGGCTTTT GCACCTCCTA 60
TCCCCAGGGT CCGCCGAGAG CCACGATTTT TTTACAGAAA ATNAGCAATA AAGAGATTTT 120
GTACTGTCAA A 131

SEQ ID NO:7784

SEQUENCE LENGTH:143

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09034

SEQUENCE DESCRIPTION:

GATCACAAAT GCATTTTTC AATCTTTACAA TATGAAATNT ATTTATAACT TTAGTTCCAA 60
AATGGACTCT GTAATGCATT TGCCCATATT TTTAGTCGTA ATGGATGTGA TTATATTNA 120
TTGAATTAAA AATCTAACAG AAA 143

SEQ ID NO:7785

SEQUENCE LENGTH:345

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09035

SEQUENCE DESCRIPTION:

GATCTTAACT TTAATAAGGC AAAACAAAAG CTTGAGCTG TTGCGTGTGT AAGTCTGTTG- 60
TGTGGATGTN CGTGTNTGGT CCCAGCCCC AGACTGGATT GGAAAAGTGC ATGGTGGGGG 120
CCTCGGGGCT GTCCCCACGC TGTCCTTTG CCACAAGTCT GTGGGGCAAG AGGCTGCAAT 180
ATTCCGTCCT GGGTGTCTGG GCTGCTAACC TGGCCTGCTC AGGCTTCCCA CCCTGTGCGG 240
GGCACANCCC CAGGAAGGGA CCCTGGACAC GGCTCCCACG TCCAGGNTTA AGGTGGATGC 300
ANTTTCNGNA ACCTCCANTC TTCTGTGTAG CANCTTTAAC CNAGN 345

SEQ ID NO:7786

SEQUENCE LENGTH:167

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09036

SEQUENCE DESCRIPTION:

GATCAGCTCT CTGTGAAAAT GGACCAATCA TCAGGATGTG GGTGGGGCCA GATTAGGGAA 60
 TAGAAGCAGG CTGTCTGAGC CAACAGTGTC ANCATGCTTG GGTCCCCTTN CATGCTGTGG 120
 5 AAACCTTTGTT CTTTCGCTCT TCACANTAAA TMTTGCTGCT GCTCAAA 167

SEQ ID NO:7787

SEQUENCE LENGTH:172

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09037

SEQUENCE DESCRIPTION:

GATCCTGGAC AAAGCTTGGC TGCCGGCTTC ATTTATTCCT GCTGATGGCT GAGAAGCATC 60
 15 TGTTTTCCAT CCCACTTGCC TGTCCCAAGT TTTGTTCCAT TTTTAAAAA TTTGTTGTAA 120
 ACTGCATGTT TTATAAAATA AAAATAAAAT ATCGTTTGTA ATTTATCTCA AA 172

SEQ ID NO:7788

SEQUENCE LENGTH:255

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09038

SEQUENCE DESCRIPTION:

GATCCGCCCC CCTTNGGCCT CCCAACACAA TTTATATTTT TGANGGTCCT TTTATGTGTA 60
 25 TTGTGCCATA TGCTTCTNAC ATCAGCCCCA ATAAATGCTT TAGGCGTTTT ATATTATCTA 120
 CCACTATTAC ATGCTCTCCA TCTCCAGACT ACCCATTCTT GCCATTGCCC TCATAGGGGC 180
 CCATTCTGAT AATTCTGGAT ATGTAGAAAA ATAAATCCTG TTTATGTGTT TAAATAAATA 240
 30 GATGTATACT ATAAA 255

SEQ ID NO:7789

SEQUENCE LENGTH:256

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09039

SEQUENCE DESCRIPTION:

GATCCCTGAC AGCAAAAAGT TTCTTTTCTG AGGCTGCCAT ACTGCCACTG TCCAGGTGGA 60
 40 GACTGAGCAA AGGAAGTCCT GGGCTGTGCC AGCTCCAGA GCTTCGGAAG AAAGAGCAGC 120
 AGCTCTCTCC CTGGGAACCA TCAGAGAATT CTGTTGATGT GTTCTGTGTC TGTCTGTCAC 180
 CTGGTCACGA GCTTCTACCA CCTTGCAAT TGTCATTAT CTTTCACTCC CTGAATAAAG 240
 TATCTATGCA TATAAA 256

SEQ ID NO:7790

SEQUENCE LENGTH:123

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09040

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

GATCTCCAC CTTTCGGGCC ATGTTGCCCC CGTGAGCCAA TCCCTCACCT TCTGAGTACA 60
GAGTGTGGAC TCTGGTGCCT CCAGAGGGGC TCAGGTCACA TAAACTTTG TATATCAACG 120
AAA 123

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SEQ ID NO:7791
SEQUENCE LENGTH:310
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09041
SEQUENCE DESCRIPTION:

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GATCTAGCTT CAGAGGAAGT CTACACCCCA TTCCCTTCTG CTTACAATGT ACCCATGATA 60
TGTGTTTAGC NNNGNAATAC TGTAACAGGA CATCACATGG AAAAATCAAA GCAGCTGGCT 120
CACTGTATTT AACTGAAAAG AATGCCTACA GATTGGATAT TTAATAGGGA AAATTAAGGC 180
ACTTTAATAA CANACTTCAT TATGTGAACT TGTGGAATAT TNNCATACAN TATACCTTGT 240
ATATTANTGC CATAGTTTTT NGAAACACTA ATTTAAACAA CANATTTNAG GGGTGCTGGC 300
ATATCCCAGN 310

20

SEQ ID NO:7792
SEQUENCE LENGTH:61
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09042
SEQUENCE DESCRIPTION:

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GATCCGAAGG CTGCGGGCAG CGTNTNATCC CGTGGTTTAA TAAAGCTGCC GCGCGCTCAA 60
A 61

30

SEQ ID NO:7793
SEQUENCE LENGTH:310
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09043
SEQUENCE DESCRIPTION:

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GATCTTGGTC TTATTTGCAC AGTTTTTGCG TCTTGTTTGC TTCTTGCATC TGATTAAC TA 60
GAATATTTCT NTTTCCCCCT TTTAATTTGT GATGTCACTT GACCCCATTT ATGTGTAGGA 120
GCACTACNCC ATTGGTTTCC AATACTGCAC ACATAAGATA CATACTTGTG TGCAGAAAGT 180
ATCTTCCTCC AGGCTTGTA TACCCITCAC ATGGAAGATT AATGAGGGAA ATCTTTATAT 240
TCTGTATAAA ANCAAANGCA AATTTATATA CTAATATCAT TTGTCTAAAA NTTTAAGTNG 300
TTTTCAAATN 310

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SEQ ID NO:7794
SEQUENCE LENGTH:165
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09044
SEQUENCE DESCRIPTION:

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EP 0 679 716 A1

GATCAGTAGA AAATGGACAT CAAGTTTGAA CAGATAAATC ATGGACAGCC TTATTGTGAT 60
TGAAATGCTT GTAGGTTCTG TGCCAATTTT CCACCACTGT GTACTTTGTT GCTATTTAAA 120
ACTGTATCAA CTCTAACGGA AGAATAAATT ATTTGTGATT TTAAA 165

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SEQ ID NO:7795
SEQUENCE LENGTH:149
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS09045
SEQUENCE DESCRIPTION:
GATCCGAGAG TNAAGGAAGT TCCCTGTNTT CCCCCTCCTT TTCCACCAGT NTCGCCTCTG 60
GCCTTCTCTG GCCACTCCTG GGAGGGACTG CCTCACCACC CCTGTCCCGC TGCCAGAAAT 120
ACACCCACAA TAAAAACCTG AAAACCAAA 149

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SEQ ID NO:7796
SEQUENCE LENGTH:98
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS09046
SEQUENCE DESCRIPTION:
GATCCAACT AACTGCCCAA GCACTACACA GCTGGACCCT GNTTCACCNT AACCCGCANC 60
TAGCAATCAA TACAGCTTCA TTATCTGAGT TGCATAAA 98

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SEQ ID NO:7797
SEQUENCE LENGTH:265
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS09047
SEQUENCE DESCRIPTION:
GATCTNTAGC CTCACCCTGN TCTCCTTCCT CCTACTGGCT GCTCAGGTGC TCCTGGTGGA 60
GGGGAAAAAA AAAGTGAAGA ATGGACTTCA CAGCAAAGTG GTCTCAGNNC AAAAGGNAC 120
TCTGGGCAAC ACCCAGATTA AGCAGAAAAG CAGGCCCGGG ANCAANGGCA AGTTTGTAC 180
CAAAGACCAA GCCAACTGCA GATGGGCTGC TACTGAGCAG GAGGAGGGCA TCTTTCTCAN 240
GGGNNGGGTG CACTCAATTG GNNNN 265

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SEQ ID NO:7798
SEQUENCE LENGTH:146
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

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CLONE:HUMGS09048
SEQUENCE DESCRIPTION:
GATCTTACCC GTGCNTGGNC CCCTCCCTC ANAGCCCATG GTAACGAACC CCTAGAAAGG 60
AGAGAACGGG CGTCAGGGGT GCACAGTCCA CAGCTGAAGA GCAAGGTTN GTGGCAGCAC 120
GGNCCGGCCC CTCACCCTCT NTNCCN 146

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SEQ ID NO:7799

SEQUENCE LENGTH:317

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09049

SEQUENCE DESCRIPTION:

GATCTCCGGA CCTCGGCTGC AGAGGCCATC GCAGCTTTN AAAAGTNAAG TGGTTAATTC 60
CCATTGGTGT CTTTNCCTTAT AGCATTTTCN TCTAACCTAT AACAAAGGGG CATTACATTT 120
ANCTTTAGAA CATGTGAATA GGAGGTTTNC TCATGACTTA CCATTCCAGC TGNATGGGAA 180
AGCAAAGCAG AAAACAGTGC CCCAAATGGA AAANAGNTAC TCACACAGAN CAAAACAGTT 240
CTNGGTCTTG TCCTTGGTCT TGTCAAACCT TGCCTGATGC TCITTCTAAN GTCNAAATAT 300
GANTGATAAG CCNGGCN 317

SEQ ID NO:7800

SEQUENCE LENGTH:372

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09050

SEQUENCE DESCRIPTION:

GATCTGGTTA AGTTGTGTAG TAAAGCATTG GGAGGGTCAT TCTTGTCACA AAAGTGCCAC 60
TAAAACAGCC TCAGGAGAAT AAATGACTTG CTTTCTAAA TCTCAGGTTT ATCTGGGCTC 120
TATCATATAG ACAGGCTTCT GATAGTTTGC AACTGTAAGC AGAAACCTAC ATATAGTTAA 180
NATCCTGGNC TTTCTGGTA AACAGATTTT AANTTTCTGA TATAAANCAN GCCNCAGGAG 240
AATTGGGGA TTTNAGGTT NCNGAATAGC CTATATATGG TGCATCGGNT AGGTCNTTAT 300
TGATTTTTTG ACCCTTTTCG GCTTTACCTN ATGGGAAGAC CCNGTTCNTT TTTAAATNAT 360
CCNGGTTTTT GN 372

SEQ ID NO:7801

SEQUENCE LENGTH:356

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09051

SEQUENCE DESCRIPTION:

GATCAACCTC TGAATTTTAC ATCATGAATG TAATCACCAC TGGAGCTTCA CTGTTACTAA 60
ATTATTAATT TTTTGCCCTCC AGTGTCTAT CTCTNAGGCT GAGCATTATA AGAAAATNAC 120
CTCTGCTCCT TTTCAATTGCA GAAAATTGCC AGGGGCTTAT TTCAGAACAA CTTCCACTTA 180
CTTCCACTG GCTCTCAAC TCTCTAACTT ATAAGTNTTG TGAACCCCCA CCCAGGCAGT 240
ATCCATGAAA GCACAAGTGA CTAGTCCTAT GATGTACAAA GCCTGTATCT CTGTGATGAT 300
TTCTGTGCTC TTCCTCTTT GCAATTGCTA AATAAAGCAG ATTTTATAAA TACAAA 356

SEQ ID NO:7802

SEQUENCE LENGTH:359

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09052

SEQUENCE DESCRIPTION:

GATCAAGCAG ATTCCACGAA TCCTCGGCCC AGGTTTAAAT AAGGCAGGAA AGTCCCTTC 60
 CCTGCTCACA CACAACGAAA ACATGGTGGC CAAAGTGGAT GAGGTGAAGT CCACAATCAA 120
 GTTCCAAATG AAGAAGGTGT TATGTCTGGC TGTAGCTGTT GGTACAGTGA AGATGCCNGA 180
 CGATGAGCTT GTGTATAACA TTCACCTGGC TGCAACTTC TTGGTGTCTAT TGCTCAAGAA 240
 AANCTGGCAG ANTGTCCGG GGCCTTATNT TNTCAAGAGC ACCNTGGGGN AAAGCCCCCA 300
 GNGGCCTNTT NTTAAGGGCA CCGTTTNGAN NTAAAAATTC TNGTTANCCC AGTTCNAAA 359

SEQ ID NO:7803

SEQUENCE LENGTH:134

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09053

SEQUENCE DESCRIPTION:

GATCCGGCCA GTTCATTGAA CAAACCTTTC CTTTTCTAA AGTCCAGAA GCCTTCCTGA 60
 AGGTGGAAG AGGACACGCA CGAGGAAAGA CTGTAATTAA TGTGTTTAA ATAAAAATGC 120
 AGTTTAGTGA TAAA 134

SEQ ID NO:7804

SEQUENCE LENGTH:62

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09054

SEQUENCE DESCRIPTION:

GATCTACTGA CATTAAGCAC TTTGTACAGT ACAAATAAA GTCTACATT GTTTAAAACA 60
 AA 62

SEQ ID NO:7805

SEQUENCE LENGTH:120

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09055

SEQUENCE DESCRIPTION:

GATCTGGATG ATGGTTATGT AGCTGCATAT ATTTGTCAAC ATTCCTCATG CTGTATGTTT 60
 AGATTTGTGA TTTTACTATA TGTAATTAA TTCCTTAATA AAATACTATT ATAAAAATAA 120

SEQ ID NO:7806

SEQUENCE LENGTH:209

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09056

SEQUENCE DESCRIPTION:

GATCTAATCT TCTGTGCTCT GTGTATTGGC GACGAGCAGG GTGAGAATAA GTGCGTGCCC 60
 AACAGCAATN AGAGATACTA CGGCTACACT GGGGCTTTCC GGTGCCTGGC TGAGAATGCT 120
 GGAGACGTTG CATTTGTGAA AGATGTCAC GTCTGCAGA AACTGATGC CCTCCTGGAA 180

GCCTGTGAAT TCCTCAGGAA GTAAAACCN

209

SEQ ID NO:7807

SEQUENCE LENGTH:59

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09057

SEQUENCE DESCRIPTION:

GATCTTAATT TTAATAATTA AAACCTGGNG TTTTAAAT AAAACCTTT TGTCTTAA 59

SEQ ID NO:7808

SEQUENCE LENGTH:100

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09058

SEQUENCE DESCRIPTION:

GATCAAAGTA ATTGCTTAAA TGTCTTGTGG AATNAGCTGT CACTTTATAG TTTATGTATT 60
GTGGTGGGTA CCTAATAAT AAAAAGATTT TTAATGTAAA 100

SEQ ID NO:7809

SEQUENCE LENGTH:118

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09059

SEQUENCE DESCRIPTION:

GATCTGTTTG ATGGTTACAT CAGTGTATAT TTGTCAAAAT TCCTCAAACGT GTATGCTTAA 60
GGCTTTGTGC ATTGTATTGC ATGTAAATNA TACTTCAATA AAGTACCAAT AGCATAAA 118

SEQ ID NO:7810

SEQUENCE LENGTH:183

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09060

SEQUENCE DESCRIPTION:

GATCCATAAC CTCTTTAGAG ACCGGCTCCT GGATGCAGAA TCCTATGAGG TTCTGGCCAC 60
ACCTGTTTCT TGTCCNTCAG GGGCTGACAC CCATGTNCCA TCCCTCTAAA CCCCTGTCCC 120
TAAACCTACC CCATCTNATG TAACTCTGGG GTTCCTTGGT CATTAGAGGG TTTNAGGTC 180
AAA 183

SEQ ID NO:7811

SEQUENCE LENGTH:175

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09061

SEQUENCE DESCRIPTION:

EP 0 679 716 A1

GATCTGAATG GTTTTAATAA GTGTCTGGNA ATTCCCCTGC TTGCACTTGT NTNTCTCGCC 60
 CACCACCACG TAAGACGTGC CTGCTTCCCC TTCTGCCATG ATTGTAAGTT TCCTGAGGCG 120
 TCCCCAGCCA TGCAGAACTG TGAGTCAATG AAACCTCTTT TTTTATAAA TTAAA 175

SEQ ID NO:7812
 SEQUENCE LENGTH:329
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

CLONE:HUMGS09062

SEQUENCE DESCRIPTION:

GATCCCCCAT TCCCCACCCT GGGAGATGAG GGGGTCCCCA TGTGCTTTTC CAGTTCTTCT 60
 GGAATTGGGG GACNCCGCC AAAGACTGAG CCCCCTGTNT CCTCCATCAT TTGGTTTCCT 120
 CTTGGCTTTG GGGATACTTC TAAATTTTGG GAGCTCCTCC ATCTCCAATG GCTGGGATTT 180
 GTGGCAGGGA TTCCACTCAG AACCTCTCTG GAATTTGTGC CTGATGTGCC TTCCACTGGA 240
 TTTTGGGGTT CCCAGCACCC CATGTGGATT TTGGGGGGTC CCTTTTGTGT CTCCCCCGCC 300
 ATTCAAGGAC TCCTNTCTTT TTTTCAANN 329

SEQ ID NO:7813
 SEQUENCE LENGTH:326
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

CLONE:HUMGS09063

SEQUENCE DESCRIPTION:

GATCTGCCAG CCCTACATCC GGCCGGGGCC CAGGGAGGGG AGCAAGGAGA AGGCAGGTGC 60
 GGCANAAAGC GGGCCCTGGA GGAAGAGGAG GGTGGCACGG AGGTCTGTGC CAAGAACAAG 120
 CAAAAGAAGC AGCTGAGGAA CCCCCACAAG ACCTTCGACC CCTCTTTGAA GCCAAATAT 180
 GCAAAGTNTG ACCAGTGTGG AAACCCAAAG GGCAACAGAT GTNTGTTTCA CCTGTGCCGN 240
 GGCTGCTGCA NGANGCGNGC CTTCAAAGNG ACTGCAGACT GNCCAGGTNA CGGTTTGCTT 300
 TTTAAACCA ATTTNGGAGA AGTTTN 326

SEQ ID NO:7814
 SEQUENCE LENGTH:20
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

CLONE:HUMGS09064

SEQUENCE DESCRIPTION:

GATCGGAATG NAAATCTAAA 20

SEQ ID NO:7815
 SEQUENCE LENGTH:248
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear

CLONE:HUMGS09065

SEQUENCE DESCRIPTION:

GATCAATGTT GAAAGTGTG TTATATGGCA AGTNTTTAAC ACATTCACAG TGTGTTGTTG 60

5 ATTTCAACTG TGAATTGTCT TACAGTTTTT TCAAACCTAG TTGTTTCTAT GGACACCTGC 120
 TCTGAATTGT ACATTGACTN CATTACTAAA GAACAAAAAT GTTCATTTT GTCCCAGTAA 180
 ATTGAGACTG CTTGTACACT TTCAGAAAAA TATGTGAATT TATAAAGNTT TTGTAGATAC 240
 TTGTCAAA 248

10 SEQ ID NO:7816
 SEQUENCE LENGTH:87
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09066
 SEQUENCE DESCRIPTION:
 15 GATCCCGGGA TTNACACTC CTTCTGTTTT GTTGCCGTTT ATTTTGTAC TCAAATCTCT 60
 ACATGGAGAT AATNATTTA AACCAAA 87

20 SEQ ID NO:7817
 SEQUENCE LENGTH:267
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09067
 SEQUENCE DESCRIPTION:
 25 GATCCACCTG NCCCTGACCT CCGACAGTGC TGGGATTACA GGCATAGCCA CCGTGCCTGA 60
 CCAGGGCTCT TTTAGCAAGG AAAACGTGAG GAATGAATGG CTGTTGGTGT GCAACAAATC 120
 ATACTTGCTA CATGTTGTGA ACCTGAAGTT ATTTGTTAGT CTGTATGAAG AATGTACCCC 180
 AGAGATGCAC CCTGTATCTG CCTATGTCT CTTACAATGG AGGTTCTCTC CTGTTATATT 240
 GCTGAATAAA CTATGTGAAC TGCTAAA 267

30 SEQ ID NO:7818
 SEQUENCE LENGTH:22
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 35 CLONE:HUMGS09068
 SEQUENCE DESCRIPTION:
 GATCTTCAAG CAAGAAAATA AA 22

40 SEQ ID NO:7819
 SEQUENCE LENGTH:90
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09069
 45 SEQUENCE DESCRIPTION:
 GATCATAGCT GAAAAATTAAT GATACTGTCA ATTTGAGATA GCAGAAGTTT CACACATCAA 60
 AGTAAAAGAT TTGCATATCA TTATACTAAA 90

50 SEQ ID NO:7820
 SEQUENCE LENGTH:25

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SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09070
 SEQUENCE DESCRIPTION:
 GATCATTAAA AAATTAATAA AGAAA 25

SEQ ID NO:7821
 SEQUENCE LENGTH:33
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09071
 SEQUENCE DESCRIPTION:
 GATCCCAATG AAACGAATTC TGTCTCCCTG AAA 33

SEQ ID NO:7822
 SEQUENCE LENGTH:23
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09072
 SEQUENCE DESCRIPTION:
 GATCTTCAAG CAAGTTAANT AAA 23

SEQ ID NO:7823
 SEQUENCE LENGTH:75
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09073
 SEQUENCE DESCRIPTION:
 GATCTTCATT TAGCTCCTTT ACTGGGATTT ATTGGATGCT NGTAAAAAAA TAAAATTTAC 60
 ACTGTNTATG CGAAA 75

SEQ ID NO:7824
 SEQUENCE LENGTH:284
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09074
 SEQUENCE DESCRIPTION:
 GATCAAGAAA GGGCTGTGGC TTGTATCAAA GGGTGTTTAA AAGGATTTAC TATCAGATTT 60
 GGCCTTGTTT TCTGTGATTT TGGNNAGCGT TCAAGGAAGC CAGGCTTTGG TCTGGATTGG 120
 ATGTTCTCAG GTCATGAATA TAATTCTCTG GGAACCCCCA AAGTTCTCAT CTACAAAGCA 180
 GAAGTTAATT GGAGCTGAAA ACTAATCAGT AAAGCCACAG CAGCCAGTCA TATGGGGGAG 240
 AAGAGGGGAA TGTTCATGG TTTTGTGGT TTAGAGGTGA GTTN 284

SEQ ID NO:7825
 SEQUENCE LENGTH:21

SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09075
 SEQUENCE DESCRIPTION:
 5 GATCCTTTGT TAACAAATAA A 21

SEQ ID NO:7826
 SEQUENCE LENGTH:159
 10 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09076
 SEQUENCE DESCRIPTION:
 15 GATCATATTT ATTTTAACTA TTTTITTAGA AAATTATTAA TTTTAATGT ATTTTGTAT 60
 TTA AAAAGCT TCTTCACTTG TTTCCCTAA NNNTTCANAT TGCTGCCCAA AAGTATGACT 120
 GTGGAGGAAA AAAAANTACT TNA AANTCC ACAC TTTT 159

SEQ ID NO:7827
 20 SEQUENCE LENGTH:328
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09077
 SEQUENCE DESCRIPTION:
 25 GATCTGTGCA GGGTATTAAC GTGTCAGGGC TGAGTGTCT GGGATTCTC TAGAGGCTGG 60
 CAAGAACCCAG TTGTTTGTG TTGCGGTCT GTCAGGGTTG GAAAGTCCAA GCCGTAGACC 120
 CAGTTTCCTT TCTNAGCTGA TGTCTTTGGC CAGAACCCG TGGGCTGTTA CTTNCNTNA 180
 GTTGAAGCG GTTTGCATTT ACGCCTGTAA ATGTATTCAN TCTTAANTTA TGAAGGTNT 240
 30 TTNTTGTACG CAATTCTCGG TTCTTTGANG NGATGGCAAC AAATTTTGGG NTTCTACTGT 300
 TATGTTGGGG NNCATTNGGG CCCANGNN 328

SEQ ID NO:7828
 35 SEQUENCE LENGTH:108
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09078
 SEQUENCE DESCRIPTION:
 40 GATCTCCTGA CCTCGTCATC CGCCCGTNTC GGCCTCCCAT AGTGCTGGGN TTACAGGCAT 60
 GAGCCACCAC GCCCGGCTGT TTTATTTCTT ATAAGTGTAC AGGANNNN 108

SEQ ID NO:7829
 45 SEQUENCE LENGTH:277
 SEQUENCE TYPE:nucleic acid
 TOPOLOGY:linear
 CLONE:HUMGS09079
 SEQUENCE DESCRIPTION:
 50 GATCATGAAG CCATCAGCTC CTCTGGGGCC AGCTANAGGA CAACAGAACT NTCACCAAAG 60

55

EP 0 679 716 A1

5 GACCAGACAC AGTGAGCACC ATGGGACAGT NTCGGTCAGC CAACGCAGAG GATGCTCAGG 120
AATTCANTGA TGTGGNGAGG GCCATTGAGA CCCTCATCAA GAACTTTCAC CAGTACTCCG 180
TGGAGGGTGG GNAGGNGACG CTGACCCCTT CTAAGCTACG GGACCNGGGT CACCCAGCAG 240
CTGCCCCATN TTNATGCCGA GCAACTTNTN GNCTGGN 277

10 SEQ ID NO:7830
SEQUENCE LENGTH:123
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09080
SEQUENCE DESCRIPTION:
15 GATCAGACCA AACAGTGCTG TTTCCCGGGG AGGAAACACT TTTTAATTA CCCTTTTGCA 60
GGCTCCCAACC TTTAATCTGT TTTATACCTT GCTTATTAAG TGAGCGACTT AAAATNATTG 120
AAA 123

20 SEQ ID NO:7831
SEQUENCE LENGTH:338
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09081
SEQUENCE DESCRIPTION:
25 GATCCAGAAA TGGAGACTGG ACCTTCCAGA CCCTGGTGAA TGCTGGAAAC AGTTCCTCGG 60
AGTGGAGAGG TTTACACCTG CCAAGTGGAG CACCCAAGCC TGACGAGCCC TCTNACAGTG 120
GAATGGAGAG CACGGTCTGA ATCTNCACAG AGCAAGATGC TGAGTGGAGT CGGGGGCTTC 180
GTGCTGGGCC TGCTCTTCCT TGGGGCCGGG CTGTTTCATCT ACTTCAGGAA TCAGAAAGGA 240
30 CACTCTGGAC TTCAGCCAAC AGGATTCCTG AGCTGAAGTG AAGATGNCCA CATTCAAGGA 300
AGAACCCTCT GCCCCAGCTT TGCAGGATGA ANCACTTN 338

35 SEQ ID NO:7832
SEQUENCE LENGTH:309
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear
CLONE:HUMGS09082
SEQUENCE DESCRIPTION:
40 GATCTAATCT CAAATACAGT GCTTAGTAAG GGAACCTTG TGCTTTGGG GATTGGGGA 60
TTAAACGAA TCTTTAAAT ATATTCCAA ATATTAAAT ATAAAGGAT ATTCCTCTG 120
ATGTTAAAG TGAGTATGGG GGTGGCACAT TATATGGAA GTNCCATGTT TGCTTAANTG 180
CTATGTAAA TAACTTGGA TATTANTGA CTTTGANTGT GAAANTCATA TGCCCTTNGA 240
45 NTATGGNTGC CTTAGCTCAT TTGACANCTG GNAGANTACT TGTNTATCTN GTTTNAGACA 300
CATTGANTN 309

50 SEQ ID NO:7833
SEQUENCE LENGTH:233
SEQUENCE TYPE:nucleic acid
TOPOLOGY:linear

55

EP 0 679 716 A1

CLONE:HUMGS09083

SEQUENCE DESCRIPTION:

5 GATCCAAAGT GGCAGGAGGT CTGTGTTGTC ATGGNGAACT GGAGTTTCTC TTGTAAGAGT 60
TCCCTCATCT GAAATCATGT ATCTNTCTCA CAAATACAAG CATAAGTAGA AGATTGTG 120
AAGACATAGA ACCCTTATAA AGAATTATTA ACCTTTATAA NCATTTAANG TCTTGTNAGC 180
ACCTGGGAAT TAGTATAATN CCANTGTAA TTTTTNNAT TTACATTTG TNN 233

10 SEQ ID NO:7834

SEQUENCE LENGTH:99

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09084

15 SEQUENCE DESCRIPTION:

GATCGGAATT TNACAGAATT GGGACTGTGG AACCNNGGGT CACTTGAATT TCCTATATT 60
TGTATGGCCA AATCAGGAAC CAGACAGTCT CCATAGTCN 99

20 SEQ ID NO:7835

SEQUENCE LENGTH:288

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09085

25 SEQUENCE DESCRIPTION:

30 GATCAGCCGA TGACAGAGAT TGTAAGCCGT GTGTCGAAGC GAAACTGGNG CCGCCACGTG 60
CGGGCNCCTGG TGCTTGACCT GTGCTGTAAC GACGAGAGCG GCGAGGATGT CGNGGGGCNA 120
NAAAANCGGT ACACCATCCG CTGACCCCGT CTGCTCCTCT AGGCTGGCCC CTTGTCCACC 180
CCTCTCCACA CCCCTTCCAG CCCAGGGTTC CCATTGGCT TCTGGCAGTG GCCCAACTAG 240
CCAAGTCTGG TGTCCCTCA TNATCCCNCT TACCTGAACC CNTNTTGN 288

35 SEQ ID NO:7836

SEQUENCE LENGTH:32

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09086

SEQUENCE DESCRIPTION:

40 GATCTAAATA AAATGGACNG TTCTACCTTA AA 32

45 SEQ ID NO:7837

SEQUENCE LENGTH:152

SEQUENCE TYPE:nucleic acid

TOPOLOGY:linear

CLONE:HUMGS09087

50 SEQUENCE DESCRIPTION:

GATCTGAAAC AGATGAAAGT TATAATAGTT TACATACCTT ACCTTAAAGT ATGGTTTTTA 60
AATACTTAAG ATGTGATATG ACATATAGTG TGTATATATA TCTNATNCTA NNGGAAAAAG 120
GGACATTAAA AAAGCTTATG TTTAAAGATA AA 152

55

5 SEQ ID NO:7838
 SEQUENCE LENGTH:108
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 GATCGTTCTT CATGGGGGTA AGAAAAGCTG GTCTGGAGTT GCTGAATGTT GCATTAATTG 60
 10 TCCTGTTTGC TTGTAGTTGA ATAAAAATAG AACCTGAAT GAAGGAAA 108

15 SEQ ID NO:7839
 SEQUENCE LENGTH:30
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 TGAAAATTTA TTAACACAGT GTTTTCACCA 30

20 SEQ ID NO:7840
 SEQUENCE LENGTH:20
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 25 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 TAATACGACT CACTATAGGG 20

30 SEQ ID NO:7841
 SEQUENCE LENGTH:25
 SEQUENCE TYPE:nucleic acid
 35 STRANDEDNESS:single
 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 CTTCTTTCTG TAGCCAGGTA ACTCT 25

40 SEQ ID NO:7842
 SEQUENCE LENGTH:25
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 45 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 TTTTCGGCGC TCCCATTTAT TCCTT 25

50 SEQ ID NO:7843
 SEQUENCE LENGTH:35
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 55 SEQUENCE DESCRIPTION:
 CACGAATTCA CTATCGATTG TGGAACCTTC AGAGG- NH2 35

5 SEQ ID NO:7844
 SEQUENCE LENGTH:37
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 CTCGCTCGCC CATCCTTATA CAGGCTCAGT TTTGTCT 37

10 SEQ ID NO:7845
 SEQUENCE LENGTH:37
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 15 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 CTCGCTCGCC CATGTATAGG GACAGCATTT CTGAGAG 37

20 SEQ ID NO:7846
 SEQUENCE LENGTH:38
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 25 SEQUENCE DESCRIPTION:
 CTGGTTCGGC CCACCTCTGA AGGTTCCAGA ATCGATAG 38

30 SEQ ID NO:7847
 SEQUENCE LENGTH:22
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 35 TOPOLOGY:linear
 SEQUENCE DESCRIPTION:
 CCAGGGTTTT CCCAGTCACG AC 22

40 SEQ ID NO:7848
 SEQUENCE LENGTH:22
 SEQUENCE TYPE:nucleic acid
 STRANDEDNESS:single
 TOPOLOGY:linear
 45 SEQUENCE DESCRIPTION:
 TCACACAGGA AACAGCTATG AC 22

50 Claims

1. A purified single-stranded DNA, a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto comprising any of the base sequences listed under SEQ ID NO 1-7837 and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.

2. A DNA probe consisting of a purified single-stranded DNA , a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto comprising any of the base sequences listed under SEQ ID NO 1-7837 and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.
3. A DNA primer consisting of a purified single-stranded DNA, a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto comprising any of the base sequences listed under SEQ ID NO 1-7837 and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.
4. A purified single-stranded DNA, a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto, wherein said single-stranded DNA is complementary to a human mRNA containing any of the base sequences listed under SEQ ID NO 1-7837 (wherein T is read as U) or any portion thereof at its 3' region, and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.
5. A DNA probe consisting of a purified single-stranded DNA, a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto, wherein said single-stranded DNA is complementary to a human mRNA containing any of the base sequences listed under SEQ ID NO 1-7837 (wherein T is read as U) or any portion thereof at its 3' region, and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.
6. A DNA primer consisting of a purified single-stranded DNA, a purified single-stranded DNA complementary thereto, or a purified double-stranded DNA consisting of said single strands, containing all or a portion of a single-stranded DNA or a single-stranded DNA complementary thereto, wherein said single-stranded DNA is complementary to a human mRNA containing any of the base sequences listed under SEQ ID NO 1-7837 (wherein T is read as U) or any portion thereof at its 3' region, and hybridizing specifically to a particular site of human genomic DNA, human cDNA or human mRNA.

Fig. 1

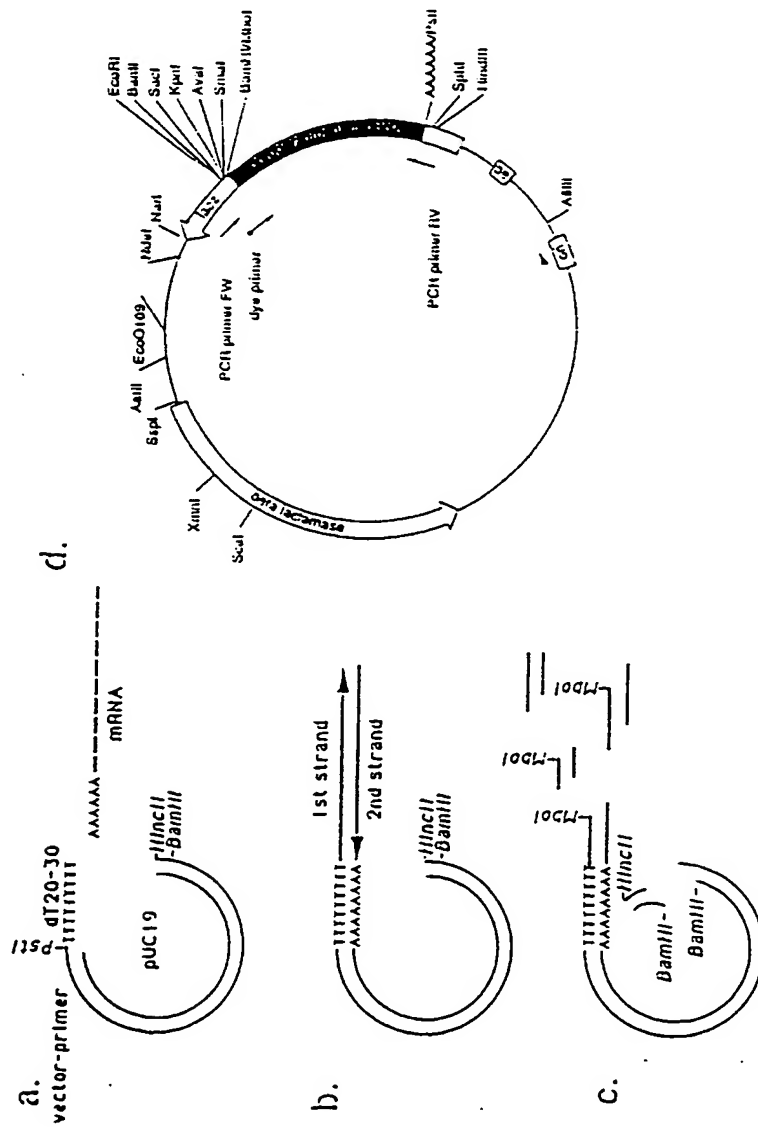


Fig. 2



Fig. 3

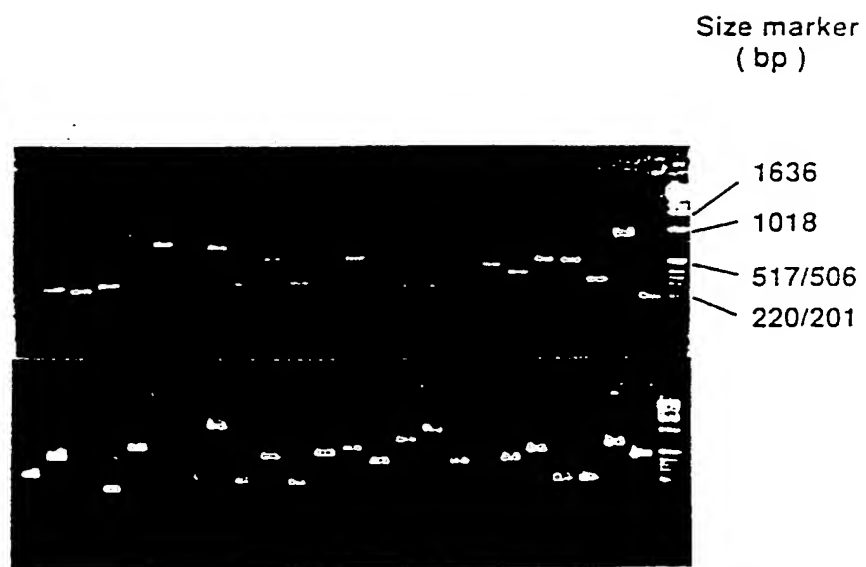


Fig. 4

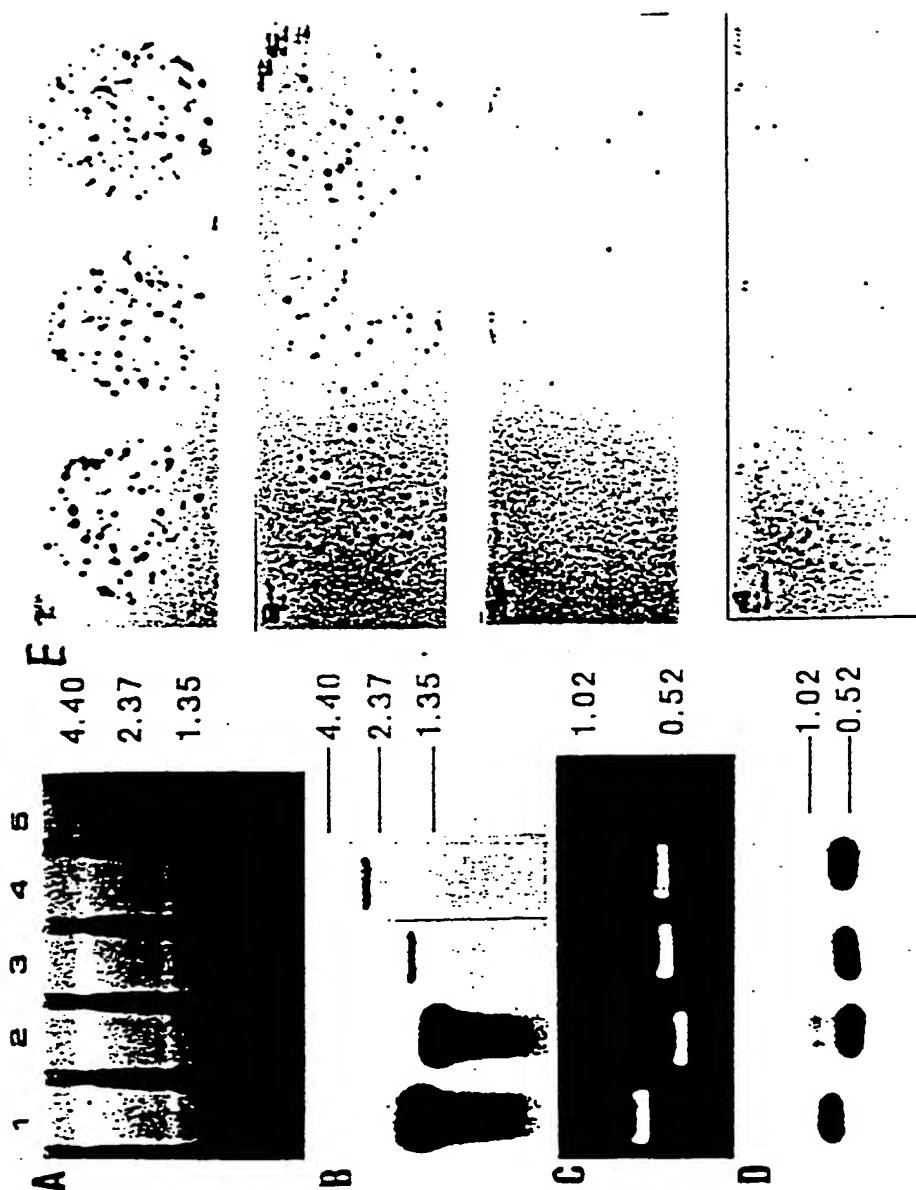


Fig. 4

F

probe No.	1	2	3	4
gene	Elongation factor 1- α	α 1-andrypsin	HnRNP core, protein A1	Inter- α -trypsin inhibitor
(a) Band intensity of Northern blot(cpm)	687	423	10	15
(b) Band intensity of control blot(cpm)	133	177	100	127
(c) Normalized signal(a)/(b) $\times 10$	52	24	1	1.2
(d) Positive signals on colony blot	307	119	7	9
(e) Relative representation	44	17	1	1.3

Fig. 5

Appearance frequencies of various cDNAs in the 3'-directed HepG2 cDNA library				
Group	Clone	Gene	A in 982 (%)	B "in 0,800 (%)" C "in 26,400 (%)"
I	a15	Elongation factor - 1A α	22 (2.2)	NT
	c321	Translationally restricted tumor protein	12 (1.2)	NT
	lb038	α -1-antitrypsin	8 (0.8)	NT
	hm01b02	Light chain of ferritin	6 (0.6)	NT
	c13a04	NADP(H) Menadione oxidoreductase	4 (0.4)	NT
	hm02d02	Ribosomal protein S11	3 (0.3)	NT
	lb042	Human RNP core protein A1	2 (0.2)	NT
	s155	unknown	1	5 (0.02)
	s159	unknown	1	4 (0.02)
	s639	unknown	1	3 (0.01)
II	s635	unknown	1	2 (0.01)
	s170	unknown	1	1 (0.004)
	s154	unknown	1	1 (0.004)
	s167	unknown	1	1 (0.004)
	s645	unknown	1	1 (0.004)
	s647	unknown	1	0 (<0.004)
	s632	unknown	1	0 (<0.004)

Fig. 6

Sequences of primers									
GS	Chl	Chromosomal position	Sette	Anti-sense	AT	IIQ	IIIE	IIIO	CO
91001070	pm2366	1	CAGAGCCCGAGTACAACTAT	AAGTTATGTGGGTCAG	48	114	115	104	110
91001076	pm2444	1	AATGGACAGTTACACTTGA	CCAGCTTCCTGACTTGAGA	48	83	81	>200	>200
91001075	pm0803	1	TGGACTGTGATGACTATCT	ACAGTACCCCTGAATGCT	48	121	124	103	107
91001047	pm1772	1	GTCACTGACGACATAGCAC	ACCACTGACGCCACACTT	50	104	104	180	>200
91001094	pm0347	1	GCCCTACACGAGGAAGCTC	TAATTTCCCACTCCCGTAAC	51	114	116	>200	>200
91001116	pm1771	1	GGGTTTCAATAGGCTGTAGACC	GCCCAATCTGTCAAACTG	49	95	95	78	107
91001181	pm0600	1	TTGCTGGATTGAACCTTTTG	GGCTGAACATTCACCTCTTG	47	97	97	200	200
91001200	pm1551	1	TTAAGAGACCCCTATGGAGACC	AATATCTTGGTATGACATAC	47	97	98	-	-
91001346	pm0882	1	TCAGTCTGCTGGAGGATG	AACCTACAGCACATGATTTG	53	120	122	>200	>200
91001446	pm1518	1	AAGGTGTACAGGATATTGCCAGA	TGCATAGCCCACTCATTT	47	130	125	>200	>200
91001464	pm1439	1	CCAAGACCTCCGCTTGAACA	TTGGGAGAGCCATAGACAG	51	100	100	>200	-
91001468	pm0427	1	TACTCAGTGGAAAGATAAAC	CAGTGGACCACATTTCTTA	40	98	98	-	-
91001521	pm2785	1	CCCAATCAAAATTGTAATG	TTTGAATCAGAGACATGAAGTT	43	102,175	100	>200	>200
91001554	pm1291	1	CCAGAGAGTTCAGGAGATG	GGTACAAAGTGCATAAGACT	46	57	57	78	155
91001572	pm2005	1	CCACATGCTCGTACGACTG	AACTTTATGCGACTTCT	44	58	55	>200	>200
91001720	pm1350	2	CATGATACTCTCGGTGTA	AAACAGTAGTCCGACAT	46	84	108	-	-
91001081	pm0931	2	AGGCTGAATGTGCGCATGCT	CCGTTATGCTACATGCT	46	119	119	93	115
91001081	pm0931	2	AAAGCATACAAATACCAA	TTCATATGTTAACAGTA	40	90	90	-	-
91001090	pm0825	2	TAATGTACACGATGAATAG	TAATGTAAATGCGAGTAA	45	88	88	-	-
91001213	pm2010	2	CCAGATGGAAGGGAAGTCT	CTGGAATAGGAATGAAACAG	47	125	125	150	>200
91001268	pm0935	2	TGAGTTTGTCTCTAATAA	GGAAATATGCTTCAGTTG	47	103	103	-	-
91001438	pm2435	2	TTTGTACCTACGTAGAGTACTT	TATCGTCAGTGGCTTTATTTG	52	137	137	>200	-
91001442	pm1671	2	TTATTAGGAGATCATTTCTGTG	ATCCGTGCCACATAGTGA	45	105	108	-	-
91001453	pm1245	2	TTGCTTCCGCTCTAAGT	AGTCCCATCTTCCACATG	45	67	65	>200	>200
91001535	pm1216	2	ATCTACTCTTTGTGAAGTG	ATGTACAAATTCGGTATGTAGG	45	75	75	170	190
91000875	pm0449	3	CGACATTCACCTCTCATAT	ACTGATTTTGGTCCCATCTG	44	68	67	-	-
91001001	pm1758	3	TCTGGCTCTTGTGTGTGCA	ATGATTATTAGGAGGAA	43	68	69	-	-
91001218	pm2424	3	AAAGAAAGCACATGCGCTAA	GGCCACAGTACATAATGTC	51	115	115	-	-
91001277	pm0668	3	GTAGTCTCTCGGCTTTAGC	ATGTATAGACAAATCCAAAG	42	90	90	-	-
91001219	pm1729	3	GGTCTGTTTATTTCACAT	AAGGATTTGATTCTACAT	43	77	77	-	-
91001306	pm1022	3	GATCCTGTGTGTGTGTGTCAGTC	AACCAAGAGGATGTTTCAGA	43	75	75	155	>200
91001418	pm2209	3	ACCCAGTCCCAATCCAGT	CTGCCAAATACAGGAAATCAT	46	83	83	160	140
91001466	pm2455	3	ATCTAGTGGCTGTAGTAT	ACACTCCCAAGCCCTTACT	55	105	105	113	>200
91000271	pm1252	4	GTCCTTGCTATCTGTGTTA	TTAAGAGATGAATTTATTTGGT	42	130	130	100	>200
				AAGCATTTATTTGAGGTTTAA	43	90	90	95	>200

Fig. 7

91000148	pm2756	4	GGCCAAAGTTCTCTAGTAT	GTCAGTTTATTCAGAGCA	42	62	62	62	69	1	2
91001052	pm1151	4	GTCCCACTGCACTGTTGTTAT	GTCAATATTCATCATCA	43	80	80	80	80	1	1
91001215	pm0988	4	AGAAATTAATAGCATAGGT	TAGAGTCAAGTTCCTGCTG	43	100	100	100	130	1	1
91001798	pm2367	4	ATCAAGTTTAAATGCTTCA	CATCCATCATACAGATC	43	116	116	116	180	1	1
91000983	pm0904	5	TCTCGTGAGAGGAGCAGCA	TCTAAGGAGAGAGCAGTC	49	101	102	113	200	1	1
91000698	pm1809	5	AGCAATGCCCTATCCACAG	CTAAGAGCTGAACCTTCAT	45	87	87	87	200	1	1
91001085	pm2319	5	TCACCCAGATTAATTCAGCT	GAGACATAGCAGGTAGAT	44	120	120	120	200	1	1
91001101	pm2364	5	TTACCTTACCGTGCTTTAC	AGACAATATCCAAAAGC	47	89	89	89	100	1	1
91001161	pm1160	5	ATTTGTGAGTGGTTTACTA	AGAAATGGATGCTTTATC	43	101	99	99	200	1	1
91000053	pm2720	6	AATGTCTAGTCTGCTTCA	TGCATCCTCAATGCTGCT	44	78	78	72	200	2	3
91001134	pm1154	6	CATTGAGACATGCTGCTCA	CCTGGCCCTCTCCCTGAT	53	102	104	145	200	1	1
91001157	pm1716	6	TAGCAAAACACGAGAGAG	AGGACCTGGTGTGAGTTC	48	65	65	110	200	1	1
91001523	pm1785	6	TATATCAAAATTCCAAAGTCTG	TCTAATATTCGGTCCCTATCT	46	90	200	200	200	1	1
91001523	pm0785	6	TGTGACGTTGTGCTCAGT	TTTAAATGTCATGTAAT	42	86	70	200	100	1	1
91001525	pm0278	6	GCACCTAAGCCTCCCAAGT	TTTATATCAGTCCAAAGC	49	138	138	200	200	1	1
91001582	pm2819	6	TCTGCATTGCACAAAGACAC	TTTGAGATTTATGATGATTC	43	62	62	200	45	1	1
91000074	pm0991	7	GACCTGAAGTGTGAATGAT	AACCTAGCTTATGGGATTT	45	119	119	200	200	1	2
91001145	pm0781	7	AGCCAACTCGGGGCTCAT	CCAGGACAGGTGAGTCAT	56	159	155	115	200	1	4
91001169	pm0219	7	AATCATTTGGCCGAGCTGA	AAGACAATTTATCCAGACA	45	88	83	130	130	1	1
91001207	pm1102	7	TCAGGCAGTCTGCTCAGATA	TTTGCAGGTAACTGTTTA	44	77	76	170	170	1	1
91001176	pm0956	8	AACAGTATTCGTTGTGACACTAG	TCCAATTAAGGCCAGTCTCAG	47	81	81	105	70	1	1
91001248	pm2527	8	TGCTCTTAATGGTGTCTAC	AAACCCXIAACACACTAAG	48	99	99	118	180	1	1
91000995	pm2708	8	TGTATGGATTTGGATTCG	CAMAGCAAAACAGCAGATA	44	95	95	85	85	1	1
91001055	pm0959	8	TTGCCATCAAAACACATACA	CTTGTGAGTTTGGTTCTG	43	55	55	55	55	1	1
91001157	pm0347	9	TTAAGAAATCACCCCTCATG	CACATGCTTATGGAACT	44	74	74	72	73	1	1
91001288	pm1245	10	AAGTATGTGCAAGATGTA	AAGAAACACTGCTTGTGG	45	138	139	200	200	2	3
91000278	pm2664	10	TGTGAATGCTATCTCTCT	GCATCGTTTCCATATCAGT	47	100	100	200	200	1	1
91001169	pm0800	11	ATCAAAACAAATCCAGA	ACTATATATAGTCCAACT	42	117	121	134	95	2	2
91001315	pm0445	11	GAATAGCTTGGAGATTTTAC	GGAGATCATACCTTCAGCA	46	100	100	84	95	1	1
91001352	pm1204	11	AAAGTGACCTTGATGGACAGTGG	TCGAGCCAAATATACATGCTACT	50	153	153	200	160	1	2
91001468	pm0558	11	AGGCTGTAAGGATTTTACG	CACATCATGTTGAGAGCTA	47	83	85	85	85	2	2
91001570	pm2810	11	AACCTCTAGTAAAGCATTG	TTATTAACCAATCCAGTA	37	47	47	125	53	1	1
91000273	pm0766	12	CTGTAAAGGTTTTGGAAATATGT	TTTCATTTTCTACCAATTAAT	42	75	75	145	200	3	3
91001163	pm2756	12	AGTGTATGGAAGACCTTGAG	GTTCTATGAACCGTGTAGC	48	130	130	103	200	1	2
91001193	pm1103	12	TCTCCCTATTCACAACTAGT	AATGATTCGTAGATAGCA	49	88	89	200	120	1	2
91001235	pm1103	12	CACAGCATAAAGAAATCATA	ACCTAATTTAGTTTCTCAC	46	100	100	100	100	1	1
91001274	pm1270	12	CATCATGTTAGTACAGCAGAG	CAGTTTGTCAAATGATATG	44	83	82	87	87	1	1
91001308	pm1355	12	AGATGTGAGTATTCCTCATGG	GAGACAGCAGTAAAGCAACAC	47	87	87	200	200	1	1
91001368	pm0368	12	CCAAAGTGTAGGTTACAG	TTCAATAGCCTTGGTTAC	47	95	165	95	200	1	1
91000159	pm12645	13	CTAAGATTTAATGCGATTC	AGTATGTTATGGCAGAGGA	46	104	104	200	200	1	2

Fig. 8

91001044	pm1659	13	TGTAGCGCTATCAGATCA	AGACAGACTTATGCCATCTA	44	109,200	109	>200	100	1	1
91001290	pm1731	13	GCTTCTGCTGCTGCTGGT	GCAGTTAATCATGGCTATCTCC	50	122	122	>200	180	1	1
91001362	pm0118	13	ACTGAATGGAACATAGTCT	TACATTACATGACATGTGA	40	61	61	95	103	1	1
91001366	pm0364	13	TGCTAGCTTTCCCTCCTTA	GAGCATTCGTGTTGTTCTTA	45	67	67	-	-	1	1
91001389	pm2301	13	CATGACCTGCTCAGACAA	GCCATCTTAAATGCTGACC	51	100	100	>200	-	1	1
91001492	pm0541	13	AAATGAATTAATAGCACT	ATTAGTTTACAGGAGAAT	41	72	72	-	74	1	1
91001561	pm0141	14	GTTTAAATTTGATTTGGG	CATCCACCTTACATTTCT	41	77	77	>200	180	3	4
91001564	pm2307	14	CGTTCCTAACTCTGAAATC	AATGCTCATTTATCTCAAG	42	55	55	>200	>200	1	1
91001576	pm2019	14	ATCACAATACCTTTAGTTG	AGCATACCTTATGGAGAT	39	69	69	-	-	1	1
91001579	pm2320	15	TCCCATCTCTAGTTGAAGT	TGAGCAAAAGGAACACAT	47	70	70	80	150	1	1
91000980	pm0985	15	TGGAAATGGAACCTTGCTA	ACTATGCTGCCGTAATGG	48	79	79	66	70	2	2
91001242	pm1127	16	CCCTGTTTTCATGTTC	TATTAATCTCCCATTCAT	44	105	105	103	102	2	2
91001516	pm2543	16	ACAGTCTAAATCAACGGTG	TGTGACACTCAAGTGCMAAT	45	70	70	>200	-	1	1
91001566	pm0913	16	TTGTGTGGACTATGTAAAT	TCAGTTTAAATGGGAACCCAG	41	53	53	>200	>200	1	1
91000806	pm1157	17	CTCTCATGTCTCTACAAAG	TAGAGGAGAACTGTGGTT	47	77	77	140	>200	2	3
91001015	pm2369	17	ATATCACCTTCCCATCCAT	TCAATACGCTCTCTCAGGC	50	80	80	>200	>200	1	1
91001156	pm0202	17	CAGAAATTAAGTGCAGCAAT	TCGTATCTGCATCTTTAAGT	45	103	100	>200	>200	2	2
91001173	pm2117	17	AAATCTGTGGTTAATTC	GTGATCTACGTGACATTCG	41	118	118	145	200	1	1
91001316	pm1878	17	TAAATTTGTGGAATCTCTGGA	ACACATTTGGGTTTGGTTTAAAC	47	100	100	95	97	1	1
91001356	pm0511	17	TGTGACAGCAGCAGCTTCAT	TCGTACATTTTAAATCCACC	45	128	128	-	-	1	1
91001495	pm0538	17	CATCTCACAGCAAGGAAC	ACCTAAGATCCAGAGAAAC	40	80	80	69	>200	1	1
91001522	pm2212	17	TGACTGCAATAAGGAGTTGT	GACATACCACTTTATTTCT	46	90	90	180	>200	1	1
91001078	pm0642	17	GTCTTCAGCAGATTTACAGGT	ACTTCTCTTGAGGACACA	45	68	68	160	-	1	1
91001117	pm1815	19	TGTGTCTCCAGCTTTGTAG	GTTACATGCCCTGGTACAG	49	65	65	>200	>200	1	1
91001467	pm0289	19	GGATCAGACCAACAGTGTCTG	GCAAGGTATAAACAGATTA	46	50	50	-	-	1	1
91001069	pm1608	19	GAAGCCCACTCGACCTCA	GGAGGATTTGGGAACCGGT	54	93	93	>200	>200	2	2
91001088	pm1879	20	GCCATGCTGTAAGTGTATGT	TTAAGAGCCATTAGCTAGGATA	48	140	140	-	-	1	1
91001146	pm1146	20	GCCCTTAGGATTCAGTCTC	ACCACCCAGGTCTTTCAGG	52	66	66	180	>200	1	1
91001122	pm0112	20	TGCTGGATGACTTCTACACG	TCCCTATCATGCTGCTGTT	49	59	59	59,115	59	1	1
91001128	pm0332	20	CTGCTCGGCTAGTCTGACTC	CMAATGCTAAGAGGACAT	49	135	135	153	160	1	1
91001132	pm0647	20	TCTGAATGATGATGGAAACA	ATCCTAGTCCCAACCCAGTA	48	109	109	-	-	1	1
91001156	pm1774	20	GGAGCCACATGGATGATG	AAATGACCCCTGGCACCTC	52	124	124	>200	>200	1	1
91001210	pm1235	20	AGCCATCTGTTATGTCTTA	GGAGCAATGAACTTCTAC	44	90	90	>200	>200	1	1
91001377	pm1701	20	TCCATGGTGTAGAACCCAG	CCACATCTCCACACAGGAGT	54	142	142	>200	74	1	1
91001385	pm2101	21	GTGAGCTCAATGCTACACAG	TTTATAGTCCACACAGAGT	45	130	130	180	>200	1	2
91001427	pm0648	21	CTTCTGCTATAAAGTAGAG	ACMAITGGTTCATAAATGA	39	58	58	145	>200	1	1
91000978	pm0912	22	GGTGTAGTGAACCAATTAG	AGTTGACCCATCTCTGTC	46	124	124	>200	>200	1	1
91001444	pm0911	22	GGTCTGTTCTCCCATCTGT	AAMAGCCCAAAAGTAGTCC	48	85,80	65	100	125	1	2
91001473	pm2231	22	TGAGCTGCACTTACCTGTGAGAG	AAGCAGGTGAGTGGGTTTCT	50	94	94	67	135	2	4
91001479	pm2328	22	TACAGCCCTCCCAAGTAAAC	TTTATCTGCACTCACTACAA	46	65	65	190	>200	1	3

Fig. 9

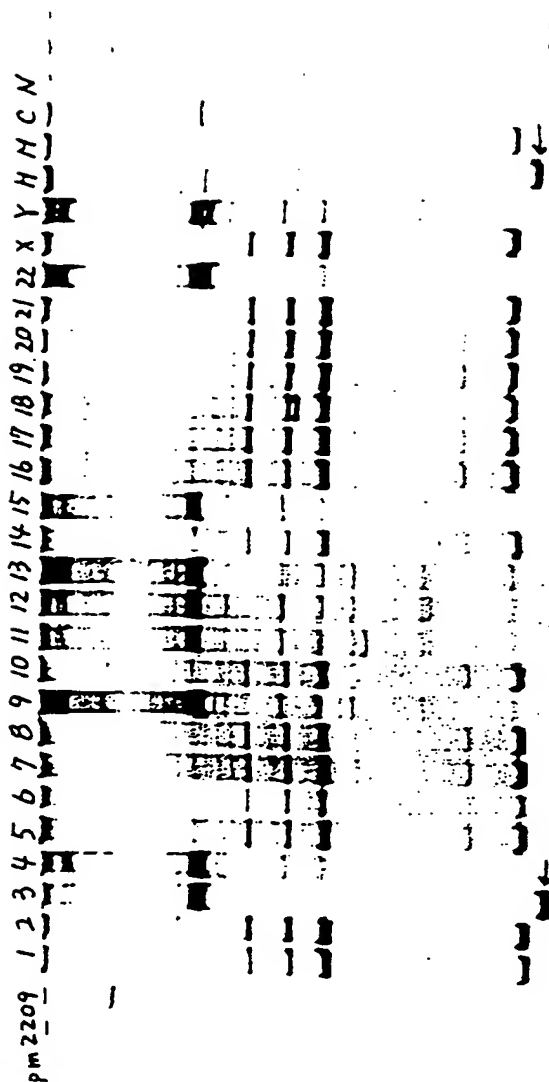
g1000999	pm1759	X	CTGCCATTAGCTGGATT	TCACCCACCACTATTAGCA	47	103	103	-	-	1	1
g1001148	pm12160	X	GGAGGGAGATATAGATTG	AAAAATCAGACAGCIGA	48	70	70	135	150	1	1
g1001161	pm0608	X	TTCATAGTGTGACCACTT	GGAGGATGAGATACAT	49	85	85	>200	77	1	1
g1001166	pm1284	X	TAATCCAGTGAATGTTGGTAA	GTAAAGGTATCTTCATCAGA	47	82	82	>200	80	2	3
g1001136	pm2289	1,18	ATCCTGCTGAATACATCTG	GGGAGACATCATCAGAC	46	70	70	68	130	1	1
g1001104	pm0113	1,2,12,13,Y	GATCCATGGGAGTGAAT	AATCAAGCTAAACCAAA	44	69	69	170	-	1	1
g1000303	pm2272	1,2,3,5,8,12,14,17,X	TTCGAAATGACATTCCTAT	TTATGTATACAAAGCACT	43	130	130	150	132	2	2
g100214	pm0214	1,2,6,X	TATCAAGTGAATGTCTAC	TTATGTATACAAAGCACT	45	93	93	110	-	1	3
g1001141	pm1461	1,3,4,5,8,16	TCCAAATGAGAGGTGTTA	AGTGCAGCAGGTTGAATG	43	96	96	100	100	1	3
g1000326	pm2785	2,20,21,22	GTCTGTCAAGCAAGATTCA	TTTATGTATGCTCCAGT	43	110	110	170	150	1	1
g1001077	pm0913	2,4,5,10,12,15,17,20,22,Y	GACCTGTGACATCTGGACT	TTATATGGTGTACACTCG	43	61	61	-	-	2	6
g1001182	pm1653	2,5,14,C	GCCTGTATTTCCACTC	ATCTCCCTTGTCCAGTTA	46	82	82	>200	82	1	1
g1000213	pm1778	2,8,12	TCTAGGAGATTCAGACAG	CAGTCAAACCAACAGGTAT	49	95	95	93	160	1	1
g1000918	pm0885	2,9,13,17,X	TGCATTTGATGCAATTC	CCGTTGTAGGTGATGAATG	49	78	78	90	>200	1	2
g1001109	pm0457	20,X	GTCAATTTGATGCAATTC	ACATTTTATTTTCAACG	37	45	45	-	-	1	2
g1000071	pm2651	3,10,15	CATGTACTAGAGGCACTTC	GCACCTACAAATCCCAAGT	50	133	133	>200	150	2	2
g1001128	pm2632	3,4,M	CAGGACTGGAGCAGGAAG	GATTTACCATTAGGAAGC	50	101	101	101	88	3	3
g1001139	pm1133	3,6	TTAGGAAATATGGTTAGACAG	ATAGTATGGGTGACACAGTA	43	90	90	>200	120	1	1
g1000377	pm2728	3,8	TGGATTGTGTTACCTGT	ACACCTCAGGAGATGTAC	47	93	93	93	>200	1	1
g1000605	pm0628	3,9,10,15	GCACACAGGCAATTCAGA	CTCTTACACCAACAGCAG	50	98	98	>200	175	2	10
g1001121	pm1234	4,6	GGATTCTATTTGCTGTCAT	GTTTATGTACGGCATTTAC	44	105	105	>200	>200	2	4
g1001132	pm1253	6,20	GCATTAAACAGGAACATA	CTGTCATGTGGCATAAACC	44	110	110	105	107	1	1
g1001441	pm0806	7,18	AGATGCTACCATTAGGATA	TTTATAGACATACAGAGGT	43	81	81	102	-	1	1
g1001157	pm0115	9,11	CCAGACTACGGCTGATGGC	CCCTTACCCAGCACTCTT	55	75,130	75	>200	>200	1	1
g1001261	pm0428	10,15,22	ACCATGTACCTGCTCTAAATA	CCATAATAGTGAAGAGTAGTTTC	48	125,155	127	125	>200	1	1
g1001154	pm2420	10,15,22	AAGAAATGTTTACTGGATT	TTATCTGAGTGGAGGAAT	42	107	107	-	-	1	1
g1000790	pm2303	10,15,22	ACTACCCCTGATATAGTT	TTCTATTTATTTGATTGTA	46	100	100	170	-	1	1
g1003114	pm2613	11,M	ATACCACTCGGCTGTACGC	GAGGAGCTCTACTGCTCTT	50	72	74	72	>200	3	20
g1003103	pm2773	12,19	GCACCAAGCAGTTCAG	TTGGGATGAGAAATAACT	46	83	83	81	-	1	7
g1001187	pm2725	12,M	GATCTCAGTCTGCGTTTAT	TACATACAAAGATGCCAACAGT	44	80	80	79	68	1	1
g1003976	pm2780	13,16	ATCTGTGTGCTGCTTCC	GTCTCTCTGATGGCIGA	46	62	60	135	180	1	1
g1001135	pm1683	14,16	AACCTGTTTTACCCCATCTT	AGGTATTTGTCCACCAGAA	48	87	87	>200	>200	1	1
g1001193	pm1748	17,20,C	TGTGTGTACCATTGAGAG	AGAACACATCAAGATCC	46	90	90	>200	90	1	1
g1000356	pm0964	17,22,Y	GAATGTATCCAGACGTAG	CTAGTTATATCTGGCTCTG	44	81	81	>200	200	1	1
g1001369	pm2217	17,C	TTATCCCAAGCAGCAAC	TCTCTCTCTACTCTCTC	49	120	120	>200	170	4	11
g1001140	pm1213	17,C	ACTTAAAGTAGCTTTGACG	TGCCCTCTGCTGTGATAATA	43	95	95	>200	95	1	1
g1001217	pm1118	18,C	CCCAGTAAAGATTATTGT	AGTAGGATGGAGGATGTA	44	92	92	-	92	1	1
g1001009	pm2824	19,20	TGCAGATGATTTCCAGAG	CTTAGGCTATCTTTTCAGC	46	75	72	160	65	1	1
g1001172	pm0887	19,22	ATCCCTCTGCTATTCACAC	GCTCGTTAGTCACTTCAC	46	110	110	130	170	2	2
		19,22	GCCTGCTCTGTTGACTT	AACCTCTGGGAACAATCAT	48	91	89	160	86	1	1

Fig. 10

91001057	pml2049	C	AGACACAAACACACGCTAT	TTTCTGATTATGACATGAC	45	75	75	101	75	1	1
91000173	pml1753	M	ATCTCTTTGTAGCCATCTGT	GTTAAGTCTGATGCCATT	42	64,100	64	110	>200	1	1
91001096	pml2236	M	GTAGAGCTCATTTGACTACC	ACAGACAGGAAATATCATA	42	108,96	110	110	112	1	1
91001166	pml0506	M	GTCCACAGCTCCAGCCTAAC	GCACATATAGAAATGCATC	46	74	74	74	>200	1	1
91001151	pml2354	M	TGCTTTGTGAGCTCTGCT	TTTAAAGCTCAATAATACATGT	44	110	110	110	106	1	1
91000279	pml2492	M1.C	GCTAGAAAGAGGGCACTCA	CTTAAAGCTGATAGCCAGGTC	46	75	75	75	75	1	1
91000253	pml2786	M1.C	CACAAACAGCAAACTTCAG	ATGGTTATTTATCAGATTG	41	93	93	82	83	2	3
91000285	pml1704	M1.C	TCCACCACAGAGAGCAGCT	ATTCATAGGGAATAGGTTTC	48	75,120	75	75	75	1	23
91000302	pml2318	M1.C	TCGAGAGGACAAATACACC	GAACAGGTTAGTCCATTGCG	48	50	58	58	58	1	1
91000313	pml1689	M1.C	CATGAGGCTACGGAAACAGG	AGAGTCCGTGGGTCTTGAAG	51	81	84	84	84	4	18
91000375	pml1442	M1.C	AMGACATCTGAGAGGAACA	GGAGGACTCGCTTGGTCTTA	49	110, >200	110	110	110	1	9
91000732	pml1452	M1.C	GCAGCAGATACCTTACACC	TGGTTCTATTCAATTCTTTC	51	102	105	102	102	3	13
91000995	pml2688	M1.C	GAAGCTCTGTGAGGAAGT	CAGACCCATCTTTATACC	47	79	79	79	79	3	4
91001016	pml2783	M1.C	ACGATATTATAGTAGTGT	TCMAACCTTTAATATAGCT	40	83	93	91	92	1	1
91001053	pml1144	M1.C	AGATGATGTGGGTACAGAGA	CCATTCTGTTCATCCAGTT	52	135	140	135	135	1	1
91001127	pml2200	M1.C	ACTGGTGATGGAGGTTACA	CCACACAGTACAGACCTTCT	47	55	55	55	55	1	1
91001167	pml1626	M1.C	GAAGCCCTTGCATCTTTTA	CTTCCCTTTGGTCTTCTGT	49	100	100	100	100	1	1
91001216	pml2100	M1.C	TAATCAGAGATTGAGTAGT	ACATGTAATTTGATAGTCT	42	110	110	110	110	1	2
91001753	pml1440	M1.C	AACTGGTTCATCAAGACTG	AGTGAATAAGCTGCCACTCC	48	120	120	120	120	1	1
91001781	pml1131	M1.C	ACTTAAACACCCACAGCAT	ACACAGCAGTCAATAGAA	47	97	97	97	97	1	1
91001375	pml0952	M1.C	AAGAGGAGTTGCCIGCTCA	ATCATGACAGTGGCAAGGA	51	89	89	89	150	1	1
91001396	pml2216	M1.C	ATCGCATGACCTATATCT	CGTTCTCTTTATTGACAT	45	108	108	108	108	1	1
91001411	pml0958	M1.C	ATGGGTTTATGAGGGTTTC	GAGACAAAGGCACTCTTA	47	80	77	80	80	1	1
91001460	pml2826	M1.C	ACATGAAATGGGATGAGGT	GGACATTTCTAGCCACAGC	51	75,55	75	75	75	1	2
91001492	pml210	M1.C	TTGTTACATTCCTTTTAGAA	CAGTCCCTCTGTACTGAGACA	48	85	85	85	85	1	2
91001608	pml2042	M1.C	GCCACACAGACATCATCT	TCCTAGAGTGTCTGGTG	51	99	99	98	98	1	1
91000450	pml204	No product	CACCAATTAAGCTGAAAGT	GAATAATCTGTGATCTA	45	87	87	-	-	1	1
91000483	pml0808	No product	CTTTGGGATTTTCTTCAT	CCCTCGGCTACTTTCTATG	43	60	60	62	-	1	2
91001254	pml1873	No product	AGCCAGCCTCTTGTATGTG	CTGGATTGATTTTCATTAG	44	87	87	-	112	1	1
91001365	pml2908	No product	TGTGATGAAATATCTGA	TTATGTAAGACACACAT	43	98	98	163	>200	1	1
91001373	pml261	No product	CAGTAGTGTCTTGAAGTG	TTTATGTAAGTGTGGTGT	41	63	63	-	150	3	3
91001556	pml849	No product	TACAGCCCTCTTAAAGTGC	TTTGAGCATCAAGGAATCT	48	82	82	-	>200	1	1
91001574	pml1284	No product	TACATCTTCAGACTCATCG	TTTCAAAAAGCTTTATCTT	40	86	86	>200	100	1	1
91001622	pml1606	No product	ATCAGAGCTCAGTCTCTGAG	ATTTCCTCTTGCATGGTC	44	57	57	67	67	2	2
91001640	pml0852	No product	GATCTGAGCCTTAAGTGGG	TTTGCAGTTCAGCTTATTC	45	54	54	-	-	1	1
			GATCTGTGTCCTTTCACA	TTTATACAGACACCATAC	36	45	45	-	-	1	1

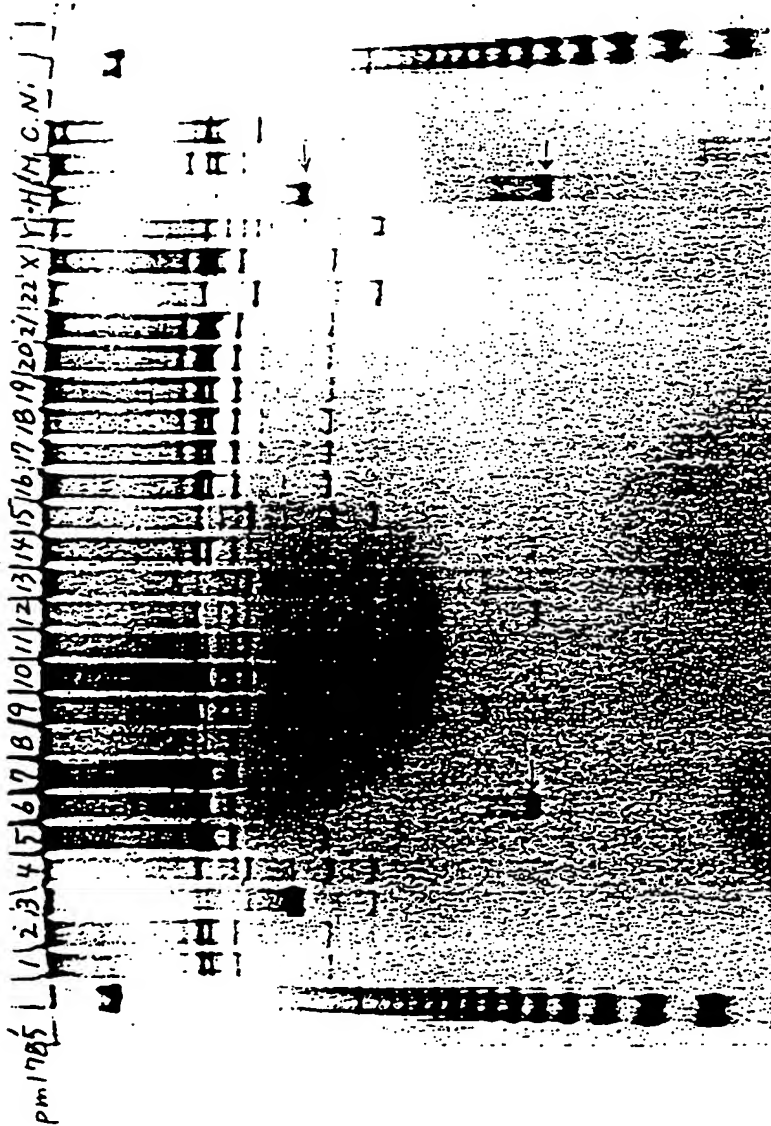
EP 0 679 716 A1

Fig. 11



EP 0 679 716 A1

Fig. 12



EP 0 679 716 A1

Fig. 13

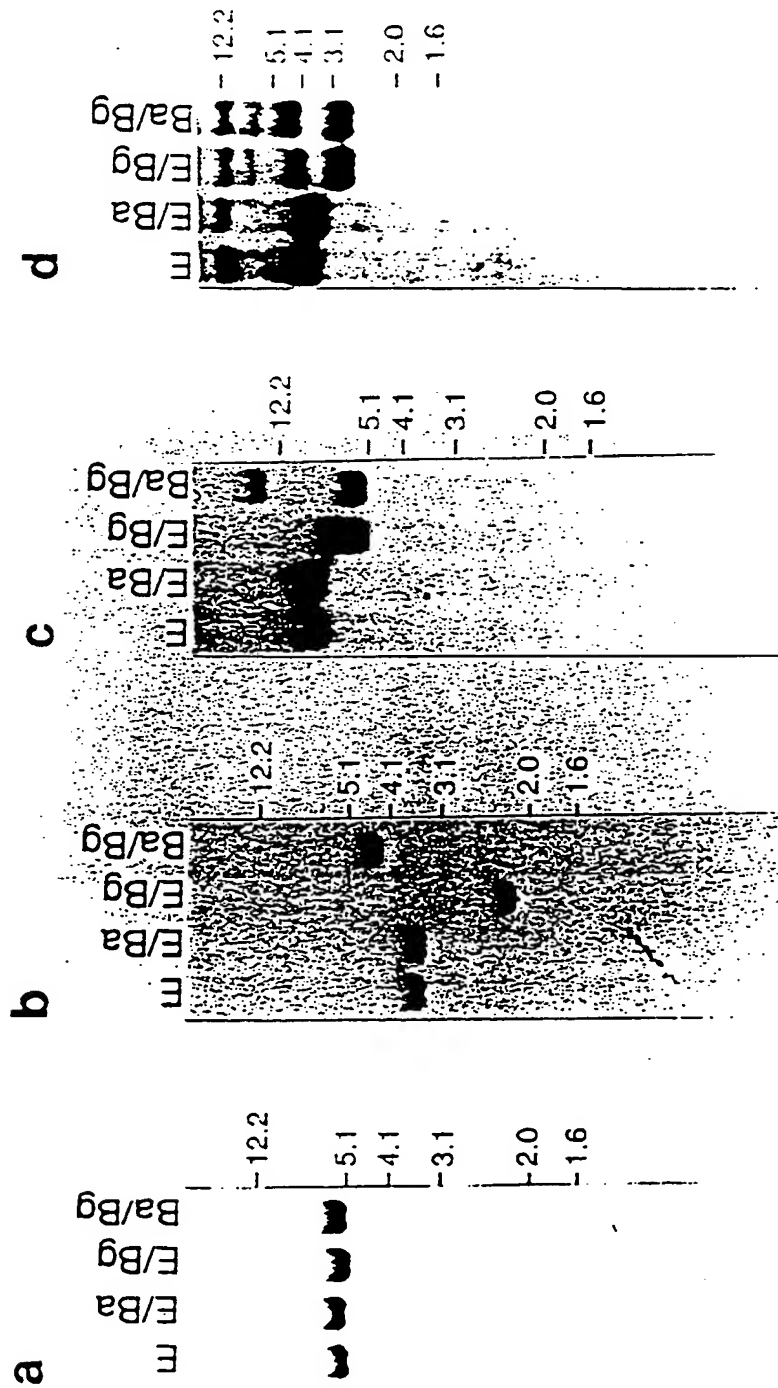


Fig. 14

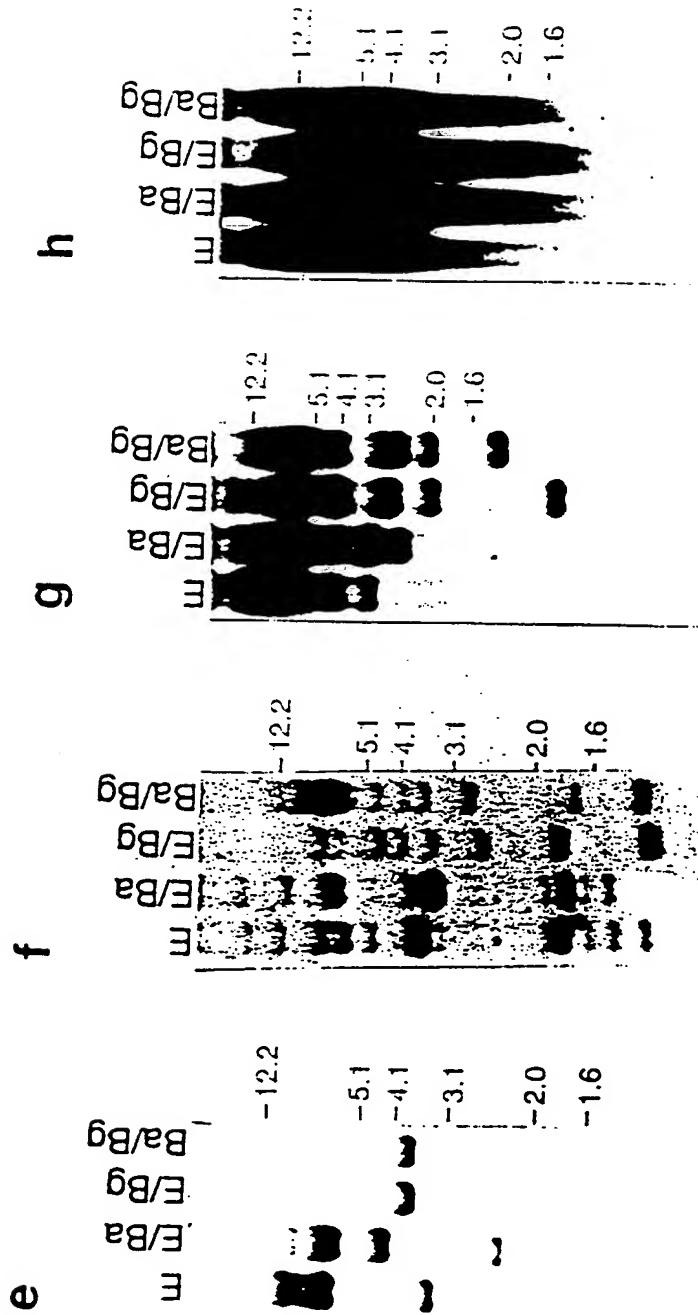


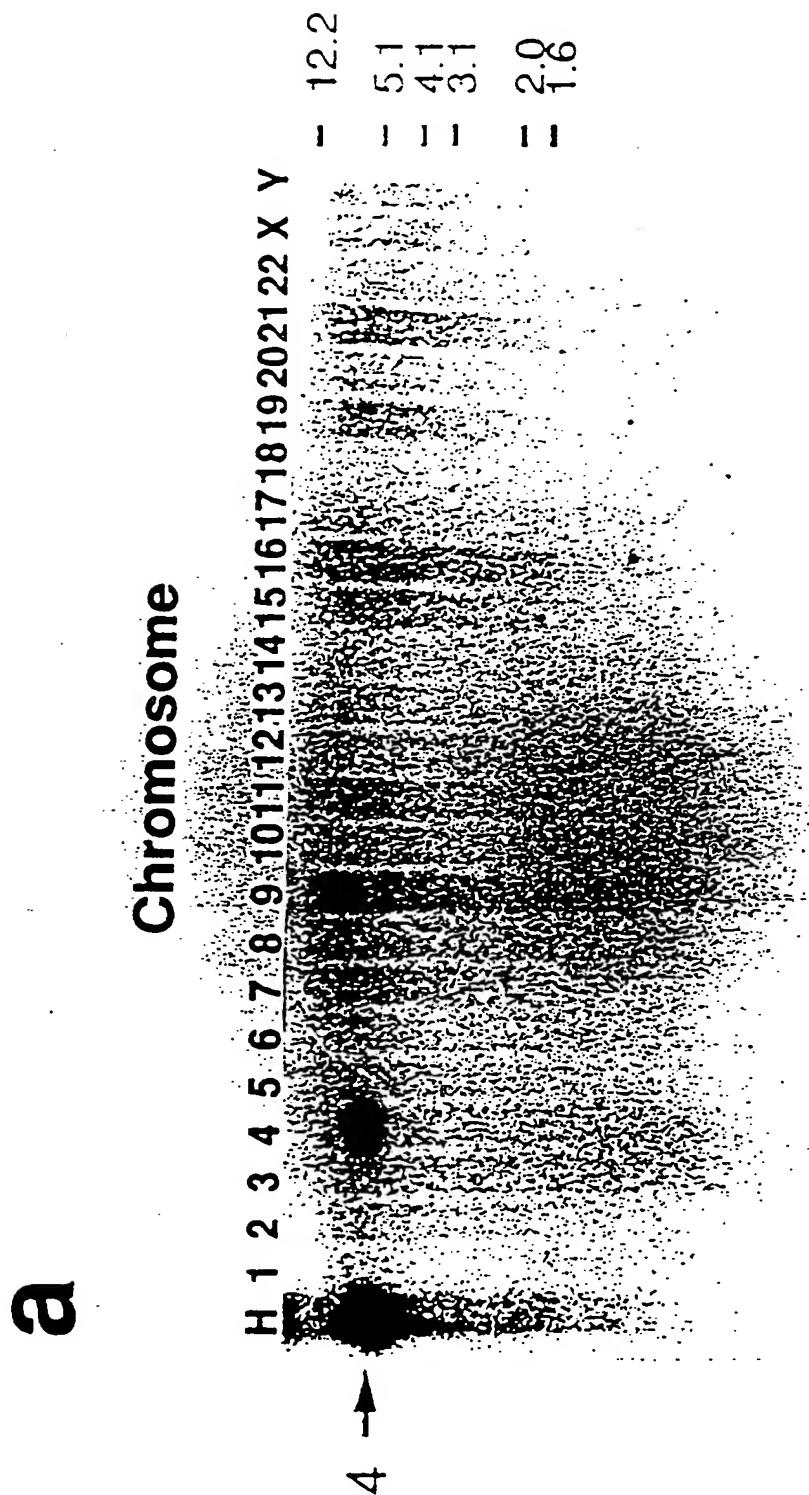
Fig. 15

Hybrid cells used for Southern hybridization

Hybrid cell	Human chromosome No.	Parent cell	Intact chromosome (%)	Translocated chromosome (%)
A9(neo-1)-4	1	A9	100 (0)	0
A9(neo-2)-1	2	A9	93 (3)	0
GM10253	3	CHO	100 (0)	0
GM10115	4	CHO	100 (0)	0
A9(neo-5)-4	5	A9	40 (0)	90
A9(neo-6)-3	6	A9	100 (60)	0
A9(neo-7)-2	7	A9	100 (89)	0
A9(neo-8)-1	8	A9	91 (82)	0
GM10611	9	CHO	79 (5)	11
A9(neo-10)-3	10	A9	94 (6)	75
A9(neo-11)-1	11	A9	24 (0)	76
GM10927A *	11	CHO	96 (21)	4
A9(neo-12)-4	12	A9	0 (0)	100
GM10868 *	12	CHO	82 (6)	0
GM10898	13	CHO	82 (0)	10
GM10479	14	3T6	76 (29)	0
A9(neo-15)-2	15	A9	9 (0)	78
GM11418 *	15	CHO	62 (0)	100
GM10567	16	A9	69 (0)	0
GM10498	17	LTMK	80 (10)	0
A9(neo-18)-5	18	A9	100 (66)	0
A9(neo-19)-1	19	A9	92 (23)	8
A9(neo-20)-3	20	A9	81 (5)	17
GM08854	21	A9	81 (24)	0
GM10027	22	CHO	93 (0)	100
GM10324	X	A9	81 (10)	0
GM06317	Y	CHW1103	91 (0)	9

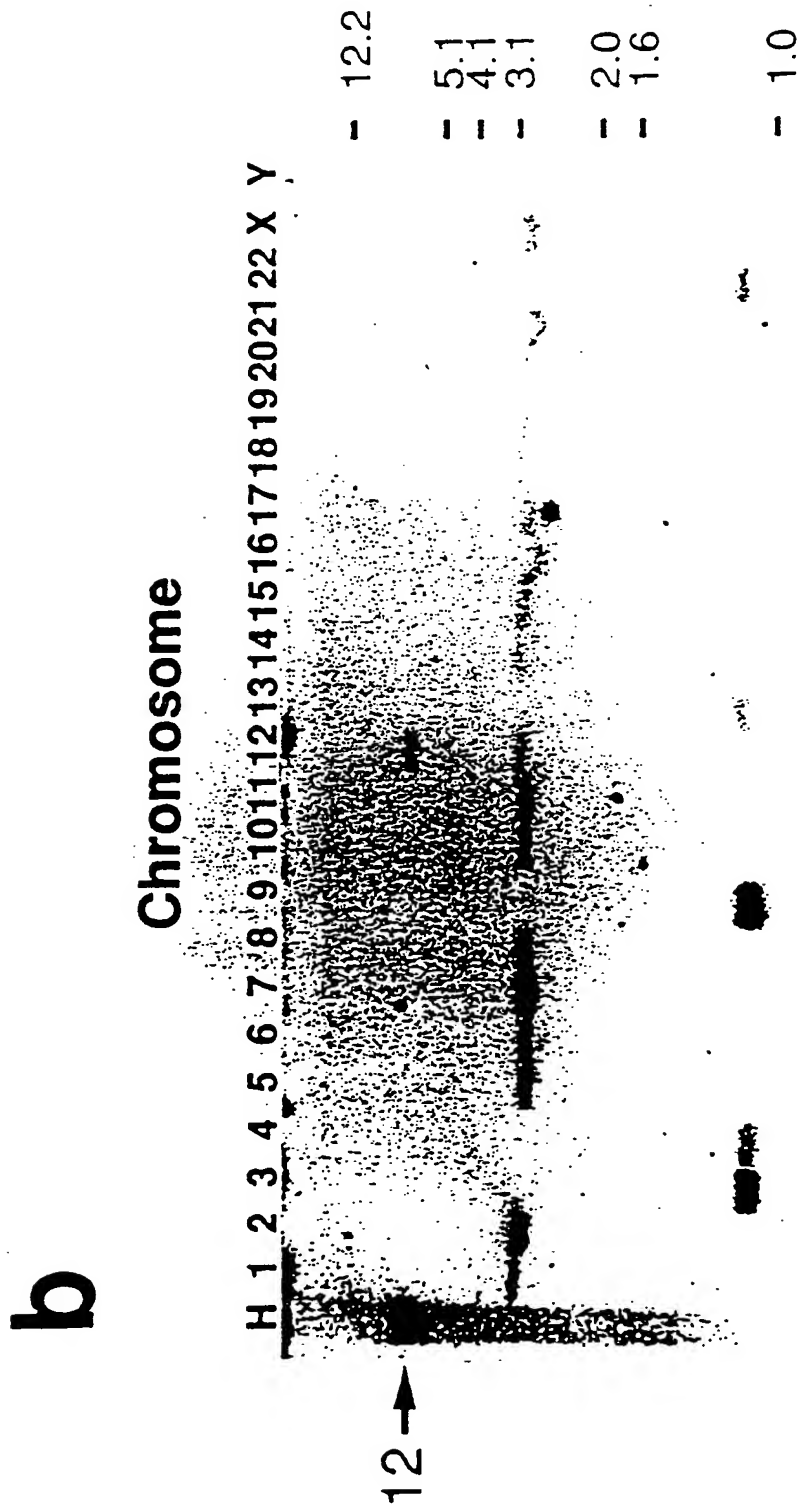
EP 0 679 716 A1

Fig. 16



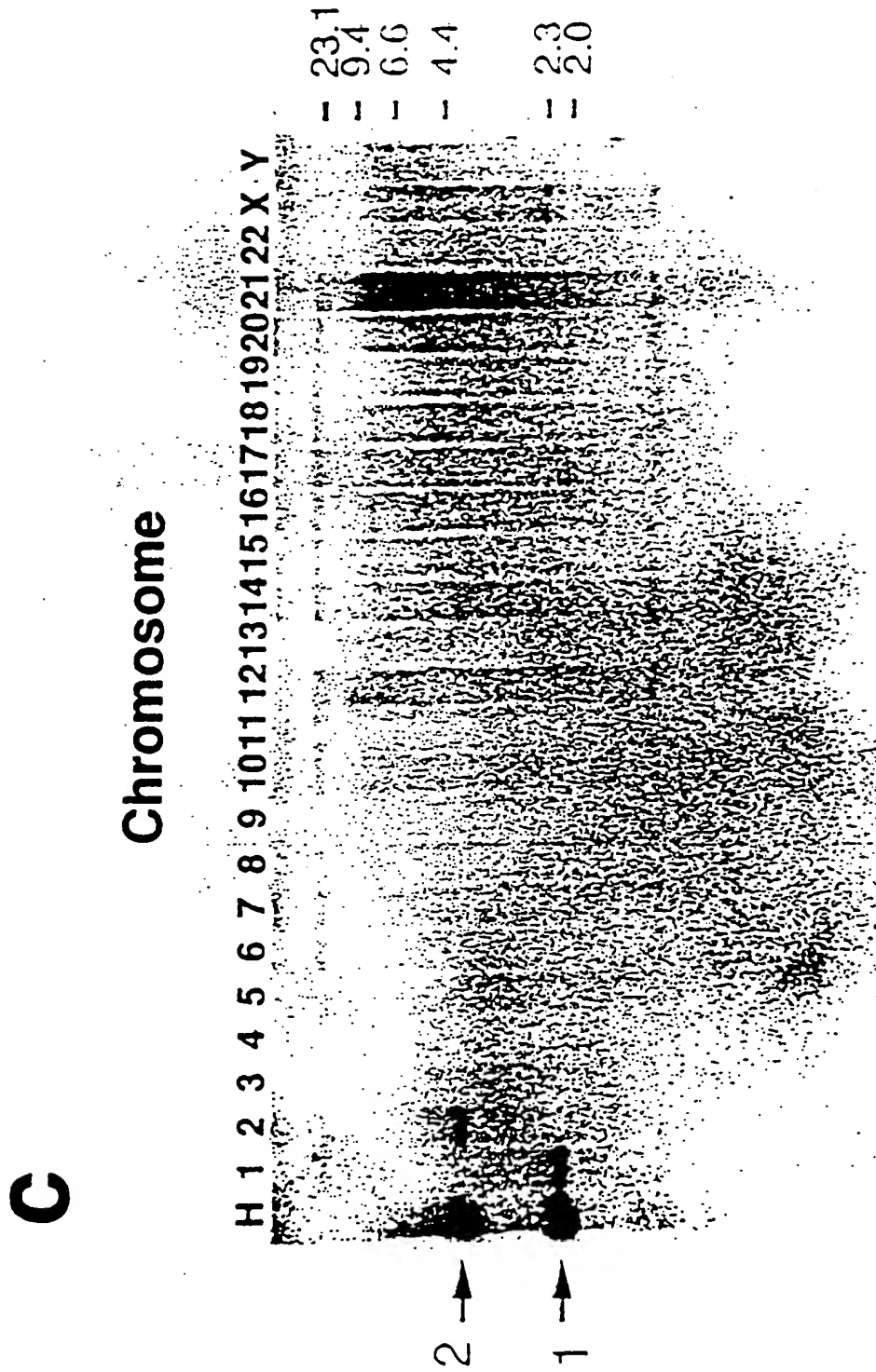
EP 0 679 716 A1

Fig. 17



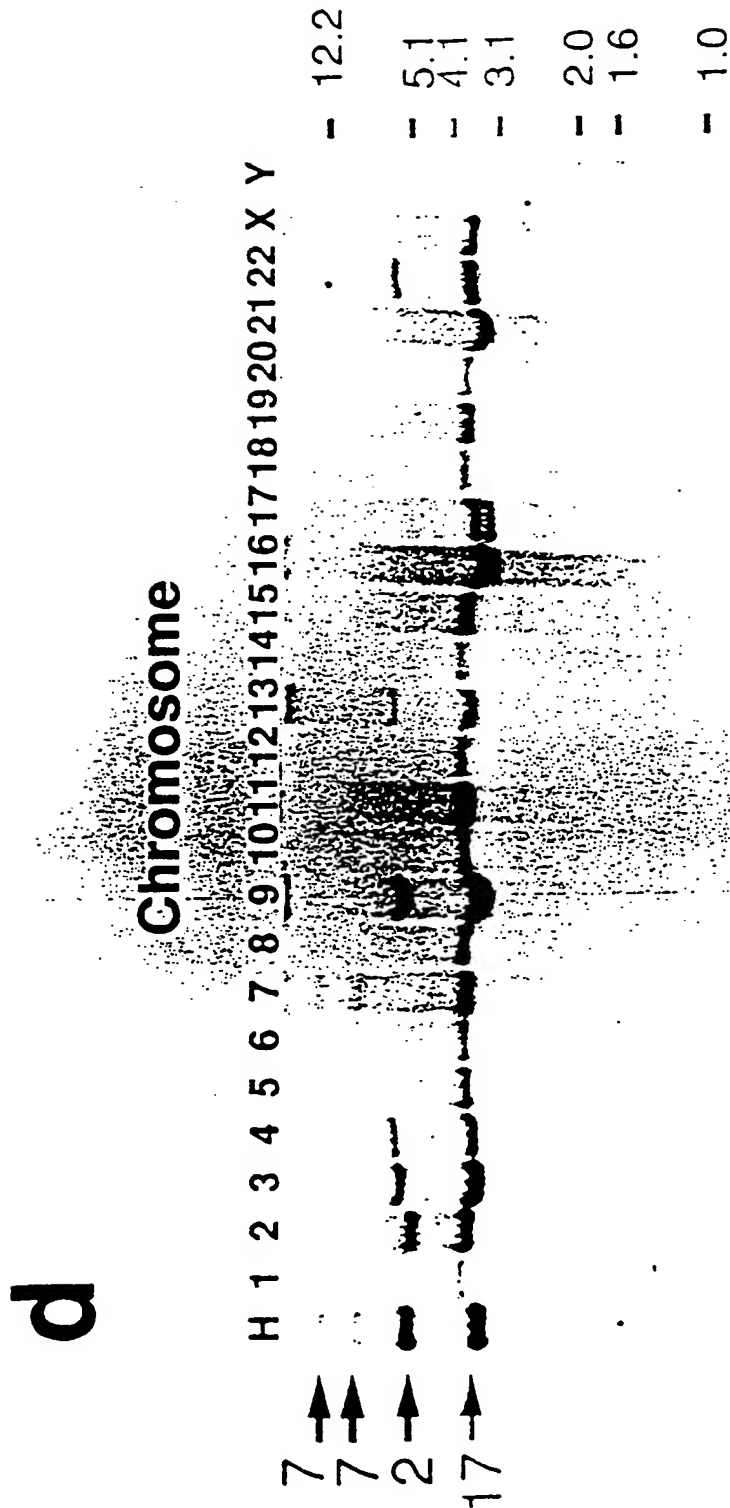
EP 0 679 716 A1

Fig. 18



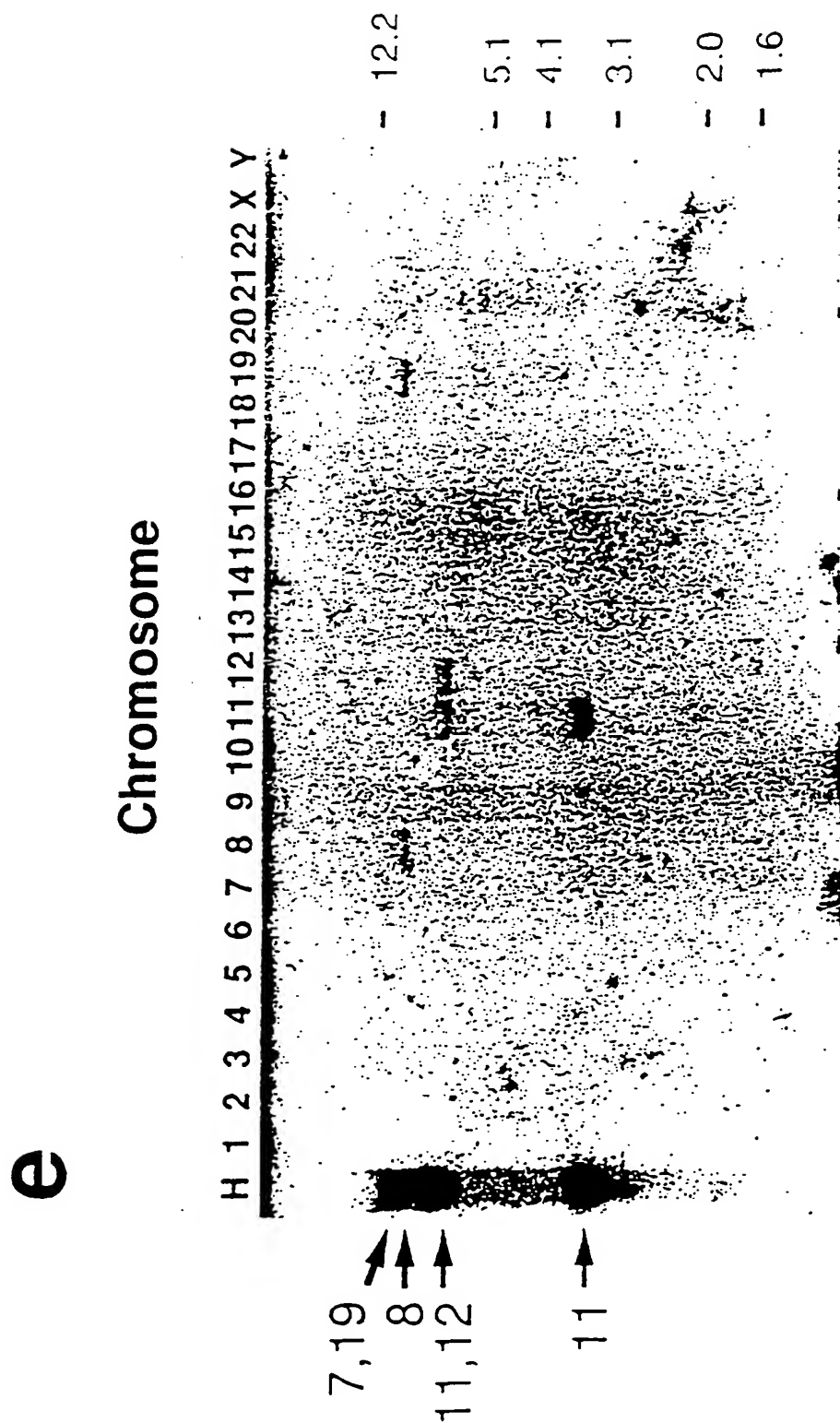
EP 0 679 716 A1

Fig. 19



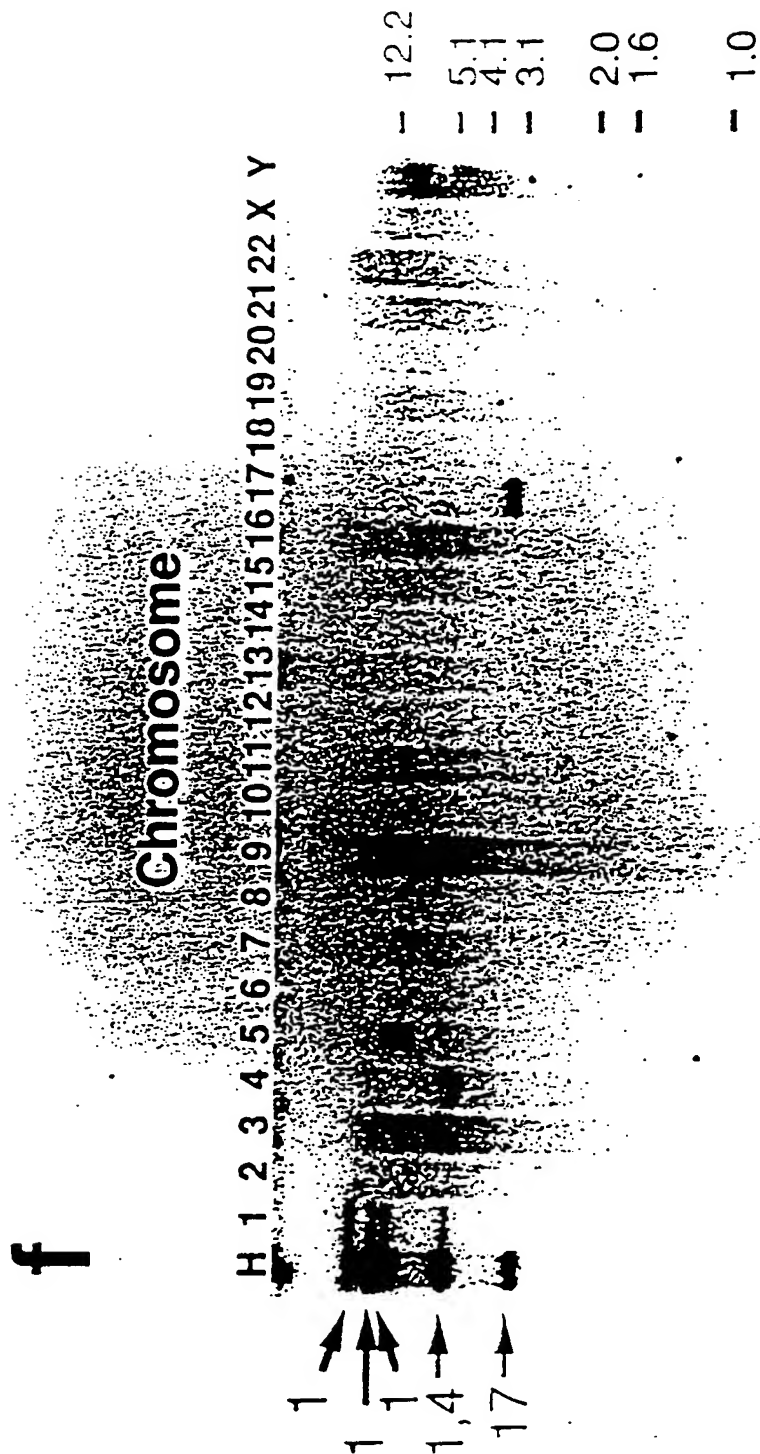
EP 0 679 716 A1

Fig. 20



EP 0 679 716 A1

Fig. 21



EP 0 679 716 A1

Fig. 22

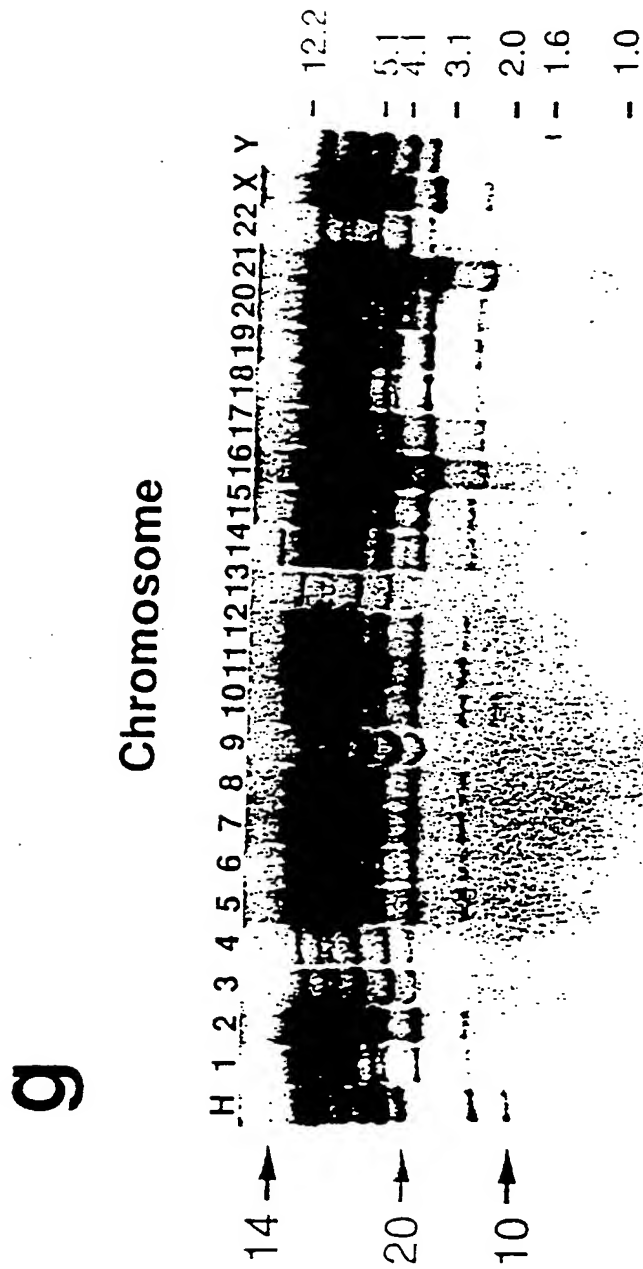


Fig. 23

Chromosomal mapping of each GS by Southern blot technique

Numbers of bands detected with human whole chromosomes		Chromosomes assigned				Background	
Clone	Sequence length \pm	2/3	3/3	3/1	3/2	Mouse	Chinese hamster
Single band group:							
c12e11	GS000075 432	1	1	1	1	9	0 0
c12e06	GS000062 540	1	1	1	1	4.15	0 0
c12g01	GS000290 212	1	1	1	1	2	1 1
c13e05	GS000117 359	1	1	1	1	11	0 0
c13e07	GS000120 355	1	1	1	1	2	0 0
c13f10	GS000206 257	1	1	1	1	14	0 0
c13h01	GS000279 133	1	1	1	1	12	0 0
c13h02	GS000222 167	1	1	1	1	6	0 0
d0g02	GS000095 397	1	1	1	1	3	0 0
d0h07	GS000134 313	1	1	1	1	11	1 1
d1b10	GS000343 153	1	1	1	1	20	0 0
hm01a12	GS000223 246	1	1	1	1	27	0 0
hm01e09	GS000423 157	0	1	1	1	1	0 0
hm01e12	junk 394	1	1	1	1	17	0 0
hm01f05	GS000066 454	1	1	1	1	19.22	0 0
hm01f10	GS000299 173	0	1	1	1	10	0 0
hm01g09	GS000053 477	1	1	1	1	6	0 0
hm01h07	GS000115 363	1	1	1	1	12	0 0
hm02a02	GS000130 544	1	1	1	1	4	0 0
hm02a04	GS000329 164	1	1	0	0	10	0 0
hm02e01	GS000292 271	1	1	1	1	16	0 0
hm02e01	GS000015 590	1	1	1	1	20	0 0
hm02e02	GS000342 156	0	1	1	1	14	0 0
hm02e05	GS000401 225	1	1	0	0	n.d.	0 0
hm02g02	GS000191 273	1	1	1	1	17	0 0
hm05a05	GS000251 219	1	1	1	1	6	2 0
hm05a10	junk 392	1	1	1	1	1	1 1
hm05e10	GS000009 606	1	1	1	1	1	0 0
km001	junk 169	1	1	1	0	n.d.	0 0
s105	GS000001 703	1	1	1	1	5	0 0
s110	GS000057 471	1	1	1	1	3	0 0
s11d11	GS000307 #175	0	0	0	1	7	0 0
s11h01	GS000253 204	1	1	1	1	3	0 0
s147	GS000050 461	1	1	1	0	2	0 0
s14e06	junk 639	1	1	1	1	1	0 0
s14g02	GS000132 323	1	1	1	1	4	0 0
s14h12	GS000271 193	1	1	1	1	4	1 1
s150	GS000142 330	1	1	1	1	17	0 0
s156	GS000002 306	1	1	1	1	2	1 1
s15b11	GS000250 221	1	1	1	1	14	0 0
s179	GS000275 196	1	1	1	1	n.d.	0 0
s246	GS000224 241	1	1	1	1	9	0 0
s247	GS000347 153	1	1	1	1	1	0 0
s270	junk 135	1	1	1	1	19	0 0

Fig. 24

Numbers of bands detected with human whole chromosomes				Chromosomes assigned		Background			
Clone	Sequence length	E	E/3 ₁	E/3 ₂	3 ₁ /3 ₂	Mouse	Chinese hamster		
s306	GS000256	205	1	1	0	1	X	0	0
s309	GS000171	305	1	1	0	1	1	0	0
s342	GS000323	165	1	1	1	1	4	3	2
s331	GS000265	207	1	1	0	1	6,15	1	1
s334	GS000165	312	1	1	1	1	1	0	0
s337	GS000276	195	1	1	1	1	17	0	0
s339	GS000295	130	1	1	1	1	n.d.	0	1
s443	GS000330	251	1	1	1	1	n.d.	0	0
s470	junk	251	1	1	1	1	17	0	0
s474	GS000192	273	1	1	1	1	5	0	0
s503	junk	312	1	1	1	1	12	0	0
s507	junk	600	1	1	1	1	1	2	1
s517	GS000334	161	1	1	1	1	14	1	1
s632	junk	387	1	1	1	1	2	0	0
s633	GS000166	311	1	1	1	1	22	2	1
s650	GS000041	644	1	1	1	1	12	1	1
tw1-04	GS000025	537	1	1	1	1	3,7	0	0
tw1-19	GS000213	255	1	1	1	1	17	0	0
tw1-32	junk	250	1	1	1	1	5	0	0
tw1-37	GS000237	235	1	1	1	1	22	0	0
tw1-42	junk	391	1	1	1	1	8	1	1
tw1-43	GS000098	178	1	1	1	1	14	0	0
tw1-96	GS000133	339	1	1	1	1	11	0	0
Two band group:									
e12f12	GS000195	277	1	2	2	2	1,	1	1
e13d02	GS000042	503	2	2	1	1	2,	0	0
hm01a06	GS000129	344	2	2	2	2	11,13	3	5
hm01a07	GS000207	269	2	2	2	2	7,	0	0
hm01d05	GS000232	243	2	2	2	1	2,	0	0
hm01e01	GS000181	292	2	2	2	2	1,2	0	0
hm02a03	GS000435	302	2	2	2	2	3,	1	1
hm02e04	GS000221	253	2	2	2	2	3,	0	0
hm02e05	GS000146	332	2	2	2	2	17,19,22	0	0
hm03f07	GS000043	503	1	1	2	1	3,	0	0
s11d06	GS000268	205	2	2	2	2	11,12	0	0
s11g12	GS000337	255	2	2	2	2	6,	0	0
s124	GS000088	404	2	2	2	2	9,	1	1
s144	GS000122	342	1	2	2	2	1,7	0	0
s14f03	GS000239	243	1	2	2	2	2,	3	2
s15e02	junk	439	2	2	1	2	6,	0	0
s16b09	junk	420	1	1	1	2	10,14	0	0
s17c09	GS000248	223	2	2	2	2	14,	0	0
s231	junk	234	2	2	2	2	11,	0	0
s254	GS000124	353	2	2	2	2	1,	3	1
s255	GS000235	239	2	2	2	2	11,	0	0
s272	junk	195	2	2	2	2	10,16	1	1

Fig. 25

Numbers of bands detected with human whole chromosomes				Chromosomes assigned				Background	
Clone	Sequence length	±	E/B ₁	E/B ₂	E/B ₃	E/B ₄		Mouse	Chinese hamster
s311	GS000092	333	1	1	2	2	16.	1	1
s313	junk	132	2	2	1	0	20.	0	0
s317	GS000100	339	0	0	1	2	14,14	1	1
s336	GS000134	337	2	2	2	2	12,14	0	0
s333	GS000139	233	2	2	2	1	22,X	0	0
s339	GS000233	137	2	1	1	2	17.	0	0
s394	GS000063	449	2	1	2	2	13,14	0	0
s396	junk	277	2	2	2	2	17.	0	1
s455	junk	452	1	2	2	1	4.	0	0
s456	GS000236	132	2	2	2	2	8,10	1	2
s465	GS000201	274	1	1	2	2	6,15	0	0
s635	junk	250	1	1	1	2	9,13	0	0
s639	GS000297	205	1	2	2	2	2,X	0	0
s656	GS000025	0590	2	2	0	2	6,11	0	0
tw1-33	junk	352	2	2	2	2	1.	0	0
tw1-39	GS000153	0321	2	2	2	2	17.	0	0
tw1-70	GS000061	441	1	1	2	1	11.	0	0
tw1-80	junk	453	2	2	1	2	9,17	2	2
tw1-87	GS000158	316	2	2	2	2	7.	0	0
Three band group									
d0h06	GS000030	417	3	3	3	1	1.	0	0
hm05b07	junk	336	2	3	3	3	5.	0	0
hm05g02	GS000209	267	2	2	2	1	3,17,19	1	1
s129	GS000107	573	3	3	3	3	n.d.	1	1
s173	GS000357	146	1	2	2	3	2.	0	0
s17a10	GS000294	131	3	3	3	3	2,13,22	1	1
s308	GS000412	638	2	2	2	3	XX	1	1
s401	GS000224	249	2	3	3	3	6,6.	0	0
s654	GS000045	491	3	3	3	3	1,22.	0	0
tw1-82	GS000208	267	3	3	3	3	13.	4	0
Four band group :									
cl2g07	GS000154	320	4	4	2	3	5, 14.	0	0
cl3a08	GS000055	508	3	3	4	4	2,7,7,17	1	2
cl3c04	GS000106	#376	4	3	3	3	n.d.	0	2
cl3c09	GS000302	195	4	2	4	4	2,17.	7	2
s136	GS000160	315	4	4	4	4	4,X	2	1
s163	GS000004	#618	4	4	4	2	4,4,8,20	3	1
s479	GS000130	293	4	4	2	2	7,8,11,11,12,19	0	0
Group with 5 or more bands									
cl2f08	GS000253	217	5	5	5	2	2,7,9,14.	2	0
hc01	junk	374	12	12	13	13	1,2,6.	22	20
hd10	junk	361	4	4	4	8	n.d.	12	6
he10	junk	173	6	2	3	3	6,3,9,19,21.	3	3
hm01c05	GS000205	176	9	7	5	5	X	9	8
hm01f04	GS000216	215	8	10	5	5	n.d.	12	12
hm01g02	junk	411	9	6	6	4	10,14,20.	14	6

Fig. 26

Numbers of bands detected with human whole chromosomes						Chromosomes assigned	Background		
Clone	Sequence length	≡	≡/3 ₁	≡/3 ₂	≡/3 ₃		Mouse	Chinese hamster	
hm02f09	GS000273	442	3	7	7	5	3,3,6,11,13,14,15,16	0	0
hm05a02	GS000096	373	5	6	4	6	2,3,17,	3	3
hm05a04	GS000236	#239	6	6	6	7	n.d.	3	5
kmb01	junk	350	5	5	5	5	13,	14	7
s11f06	GS000315	170	6	6	6	4	1,2,2,3,4,6,13,15,	0	3
s14f01	GS000407	262	12	11	10	9	1,5,9,15,	6	5
s173	GS000094	397	5	4	6	3	1,1,1,1,4,17	0	0
s255	GS000323	167	10	12	11	14	13,	9	5
s341	junk	494	9	9	3	6	n.d.	15	3
s406	GS000113	364	6	7	5	4	2,7,8,13,20,20	4	1
tw1-46	junk	593	9	10	10	10	1,1,2,2,5,11,X,	3	5
tw1-63	junk	203	3	10	10	12	3,4,	17	11
Bands no detected:									
cl3g02	GS000340	157	0	0	0	0	-	-	-
hm01a10	junk	232	0	0	0	0	-	-	-
hm02d11	GS000274	196	0	0	0	0	-	-	-
s323	GS000273	194	0	0	0	0	-	-	-
s359	GS000199	279	0	0	0	0	-	-	-
s511	junk	233	0	0	0	0	-	-	-
s645	GS000012	#734	0	0	0	0	-	-	-
s647	GS000105	360	0	0	0	0	-	-	-
s651	junk	540	0	0	0	0	-	-	-

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